

FIG. 1A

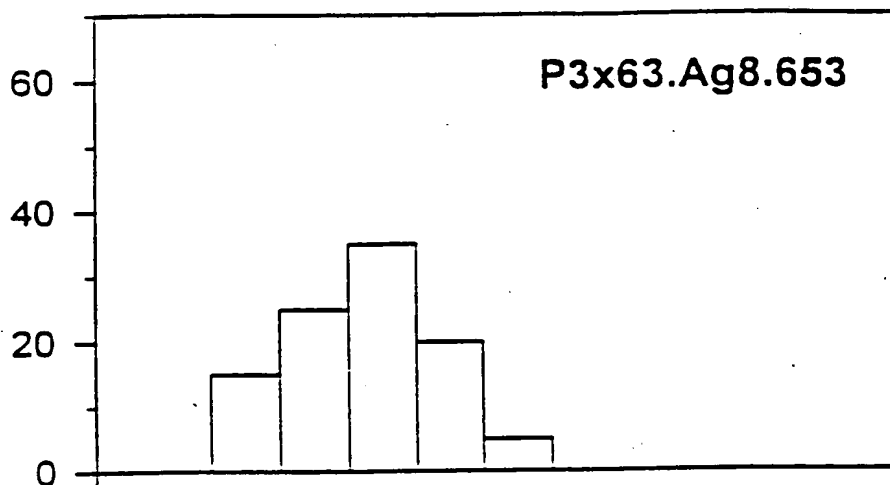


FIG. 1B

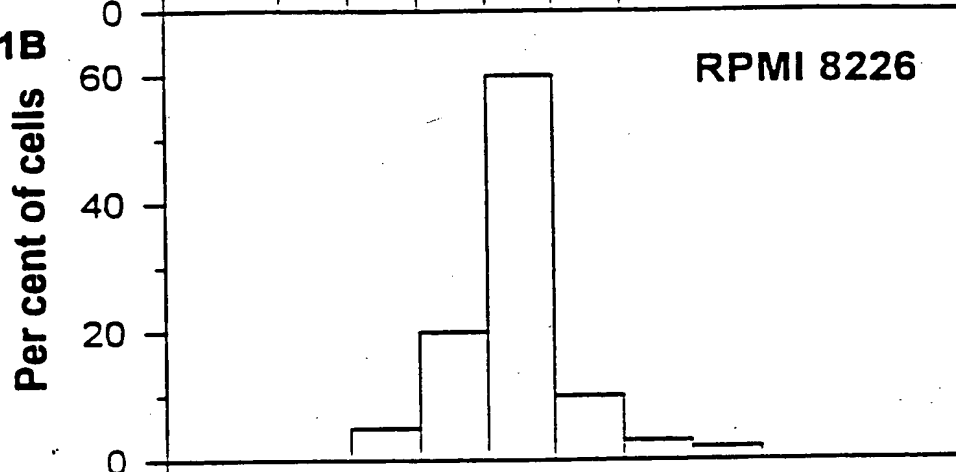


FIG. 1C

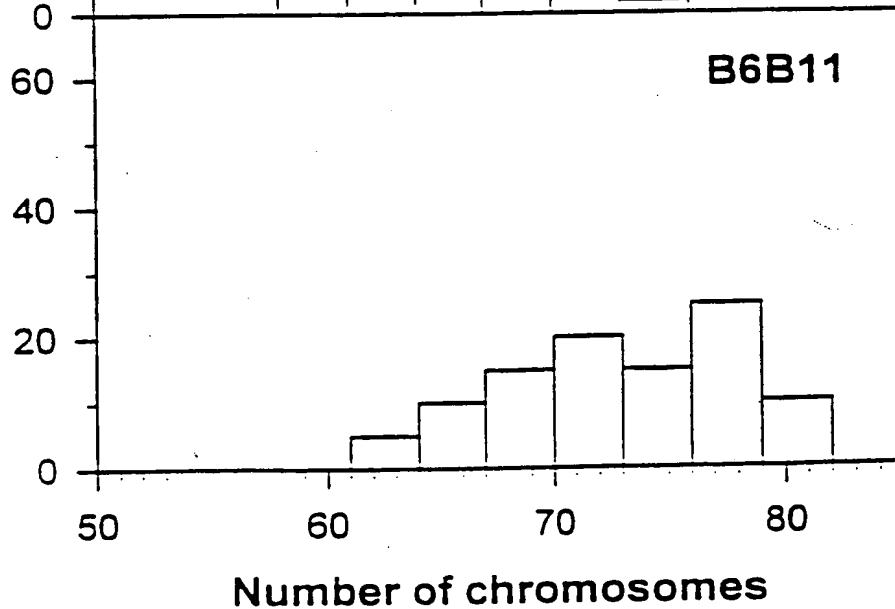
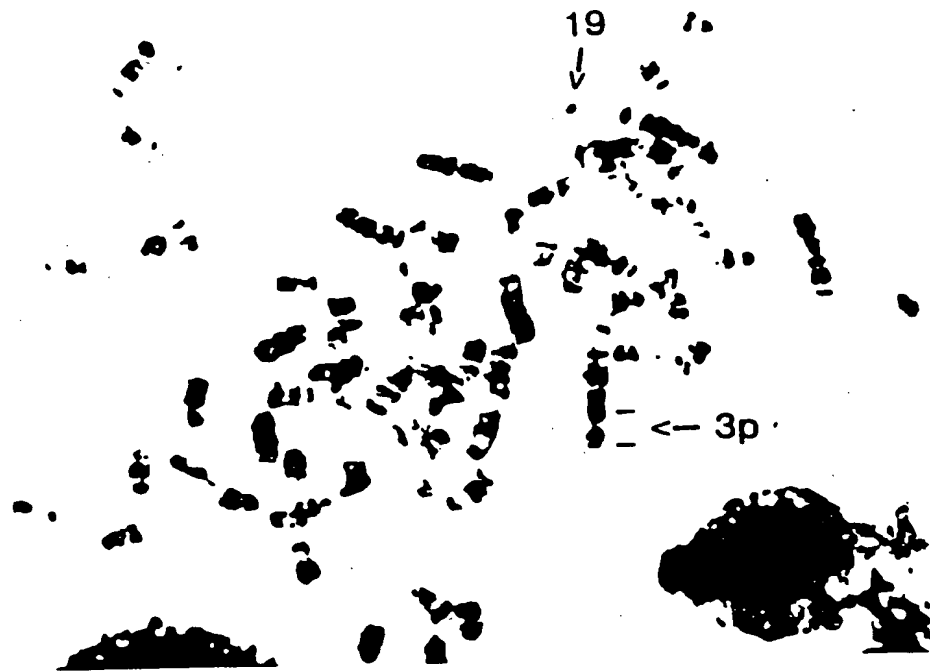


FIG. 2



008160" 8564950

FIG. 3

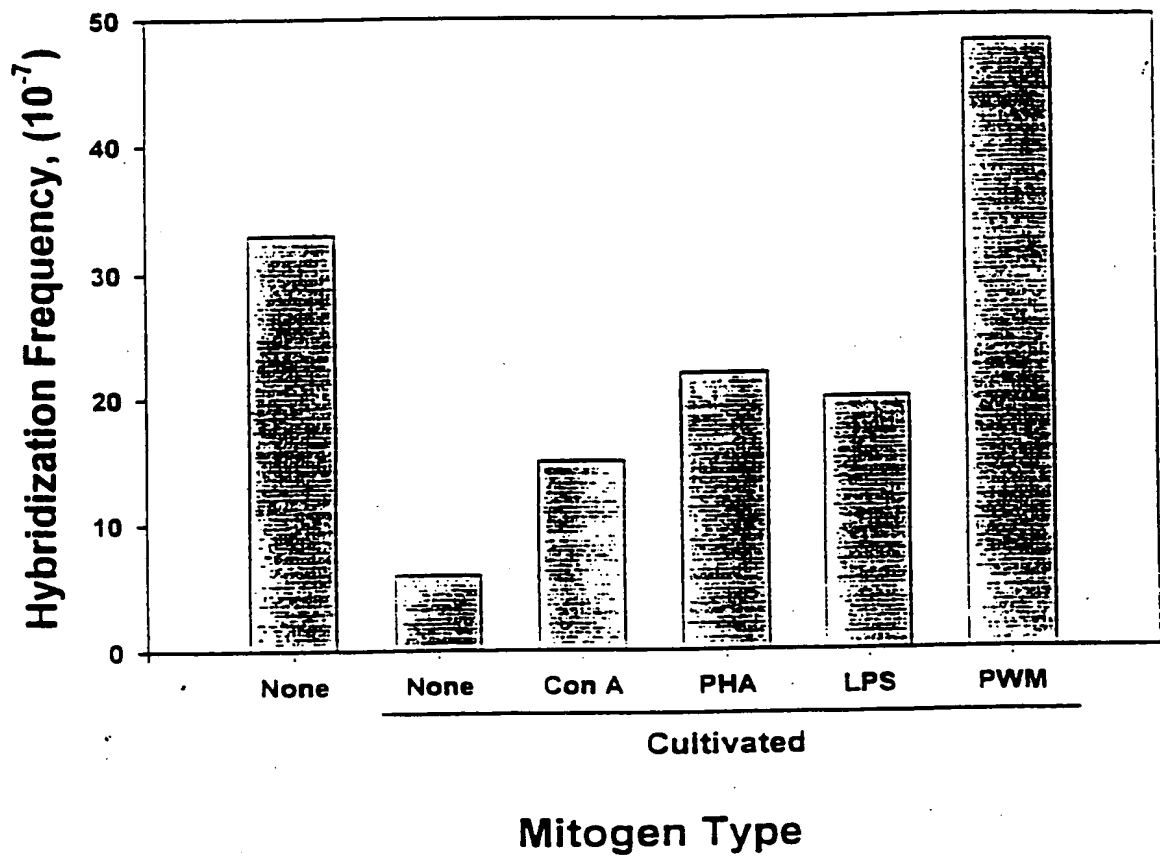


FIG. 4A

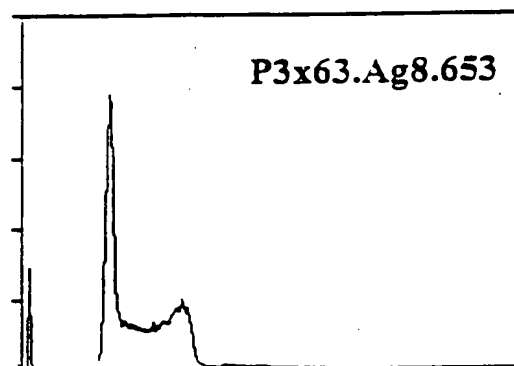


FIG. 4B

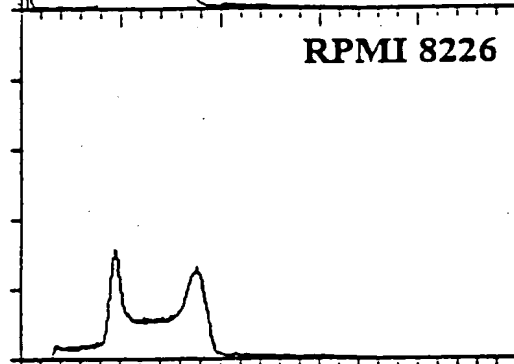


FIG. 4C

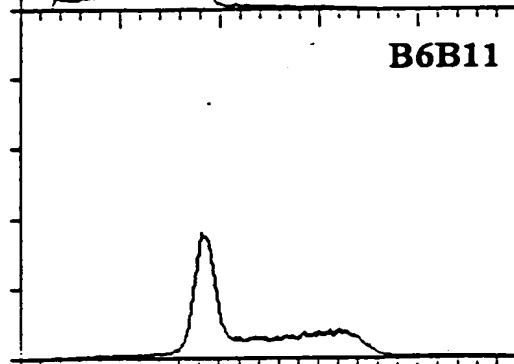
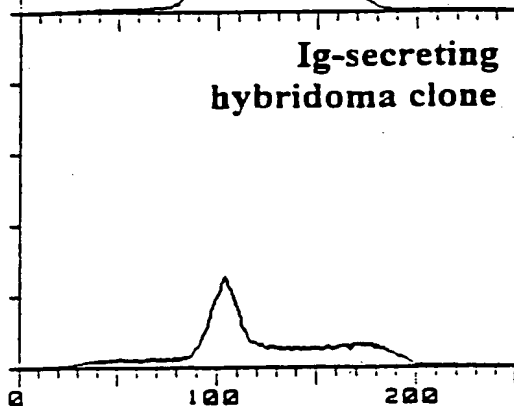


FIG. 4D



008760" 85649960

FIG. 5A

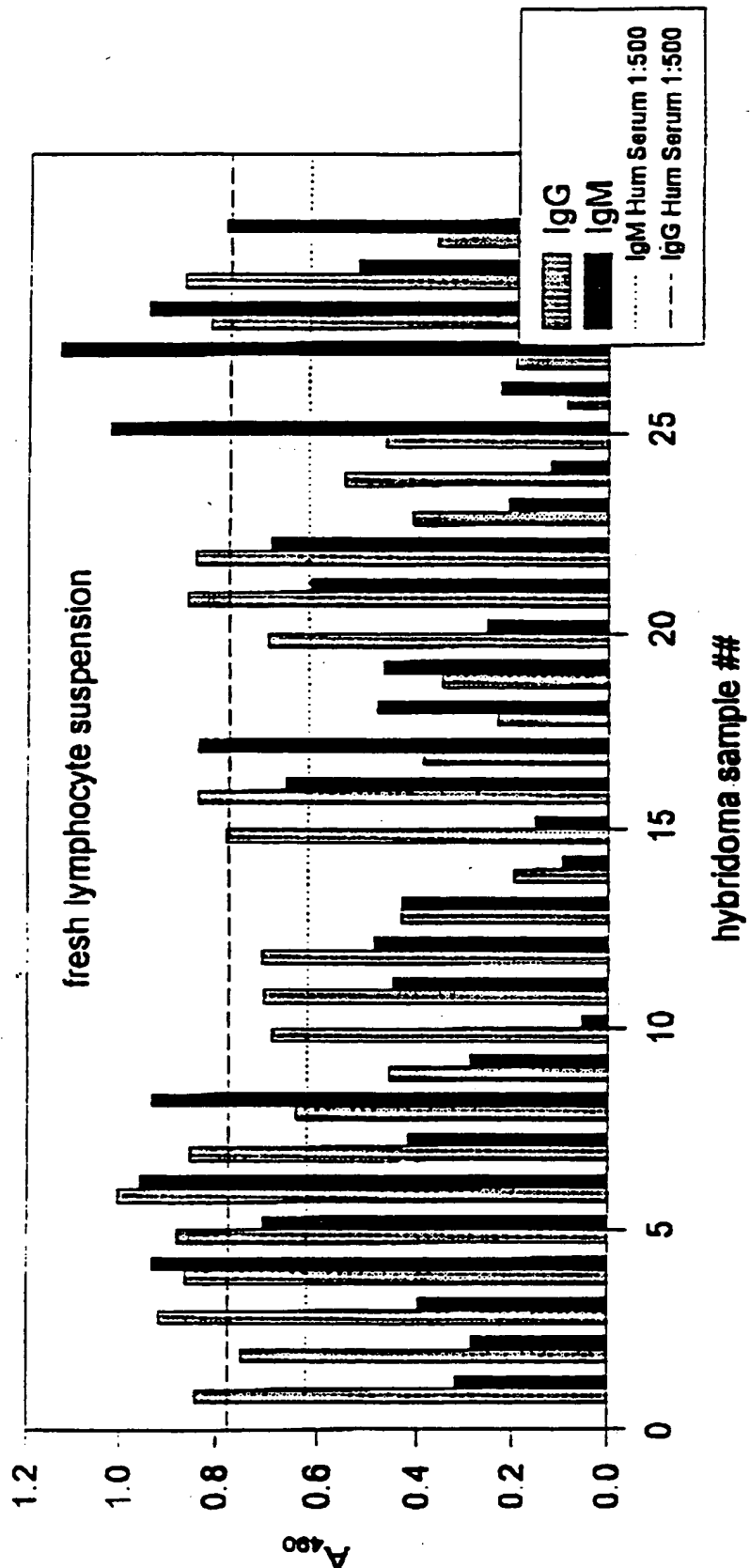


FIG. 5B

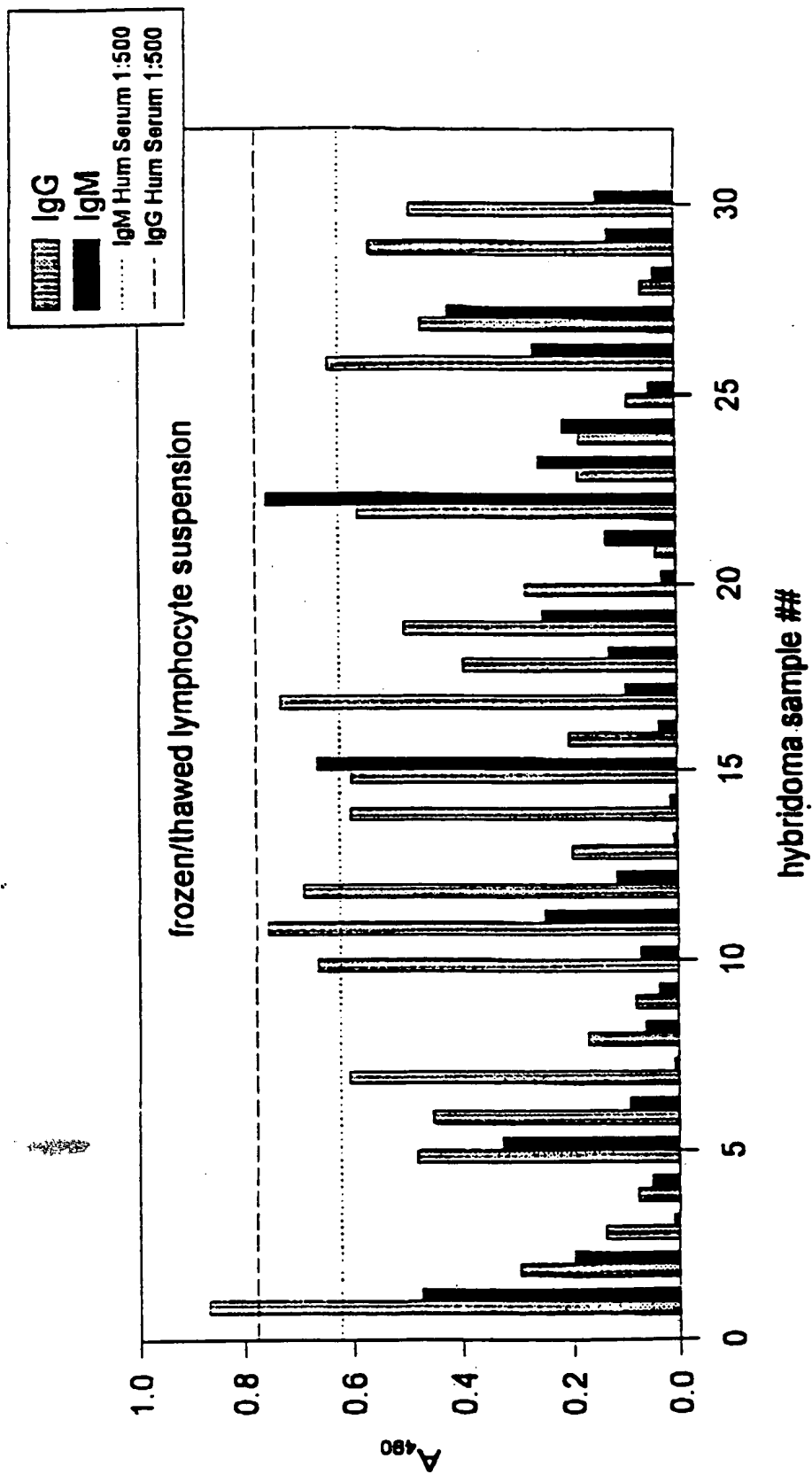


FIG. 6

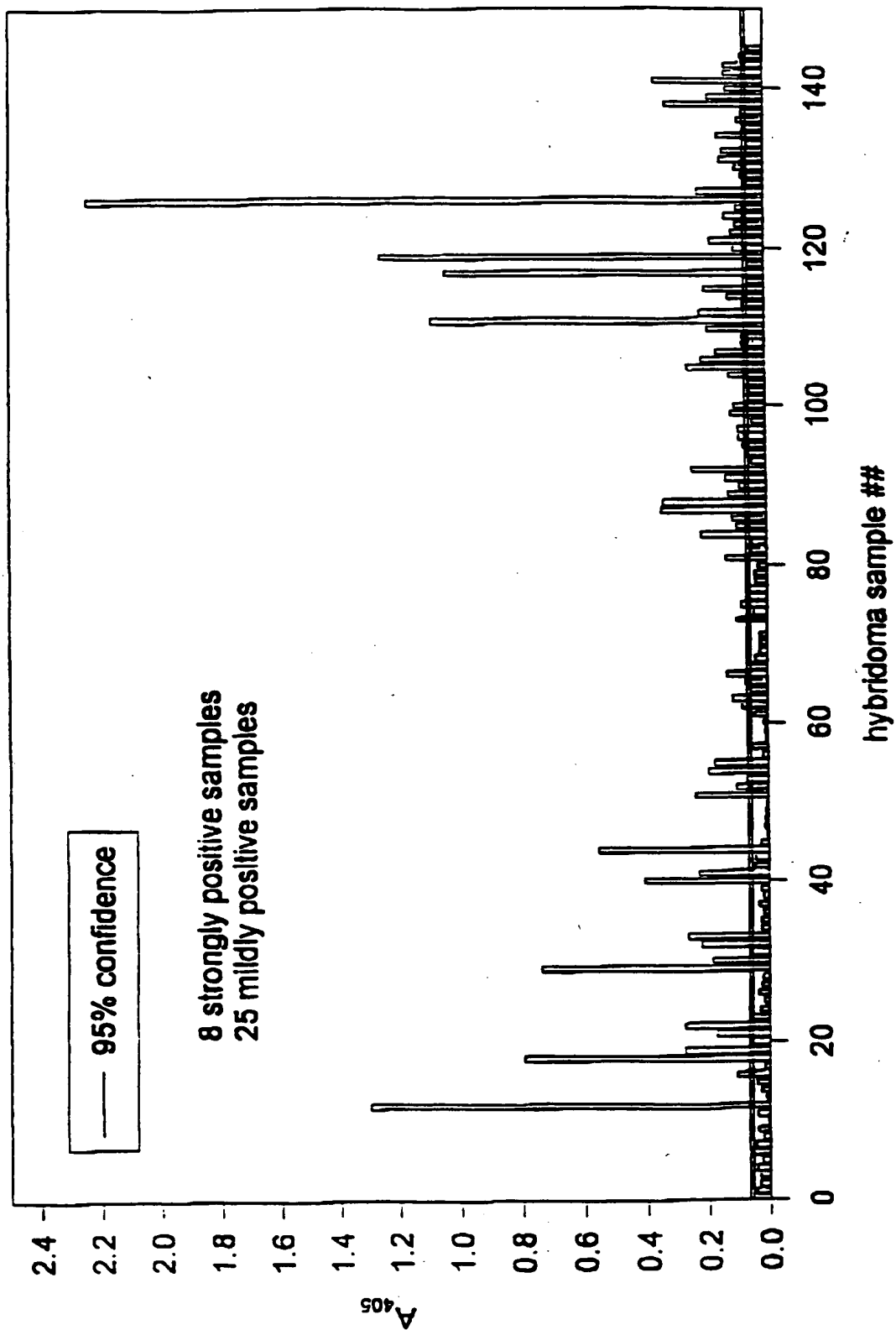


FIG. 7

27.F7

27.B1

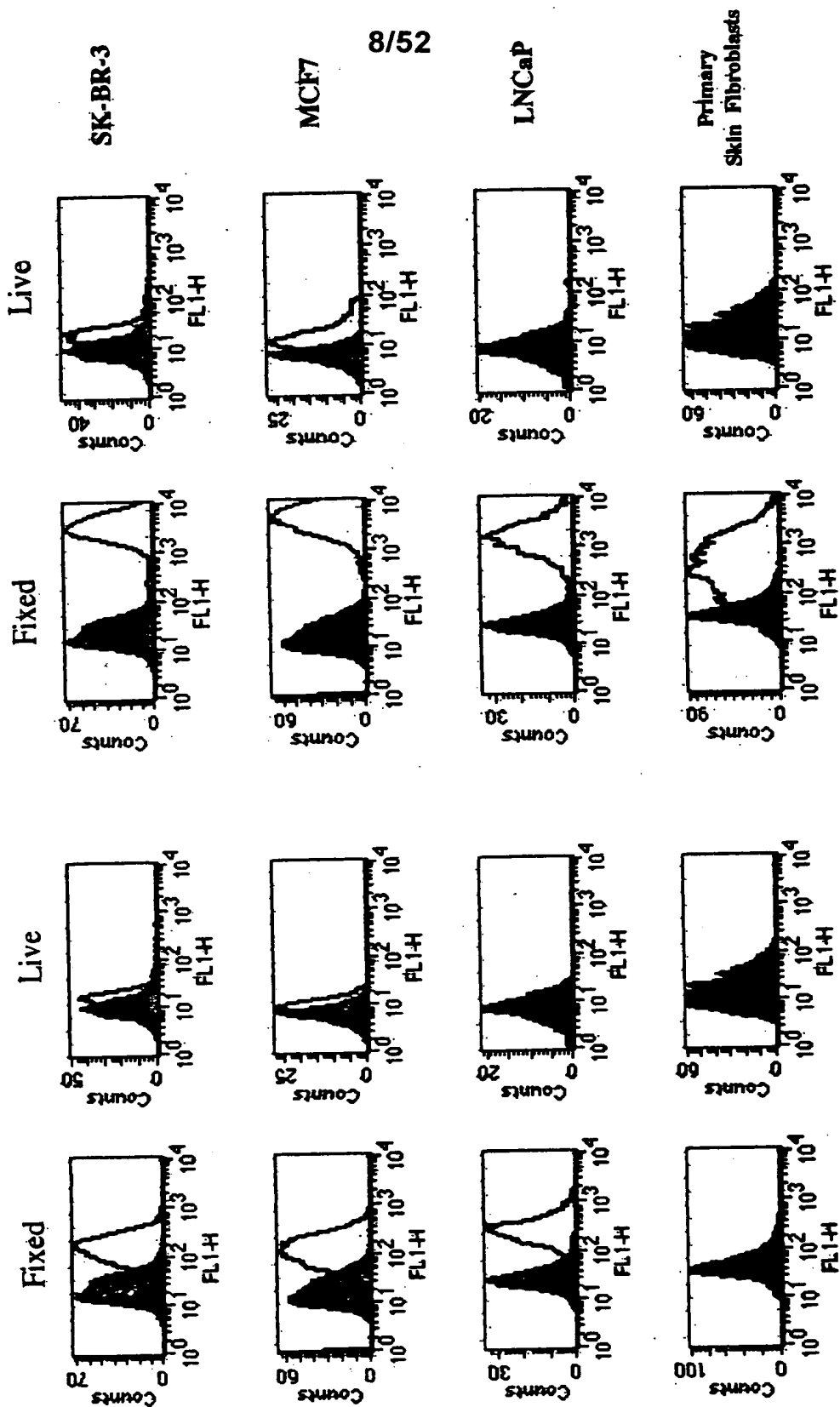




FIG. 8

# Expression of 27.F7 and 27.B1 Antigen on Different Human Cell Lines

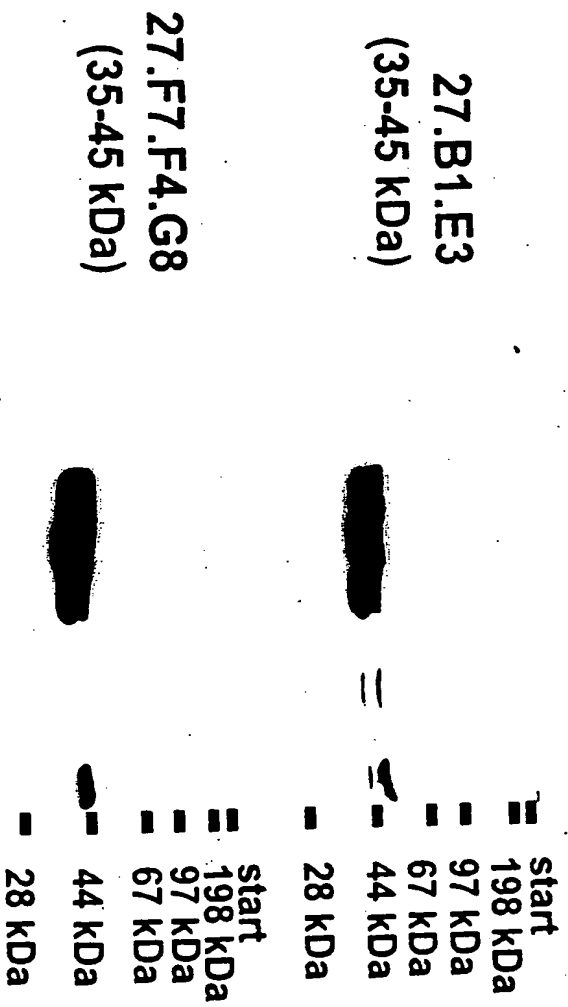
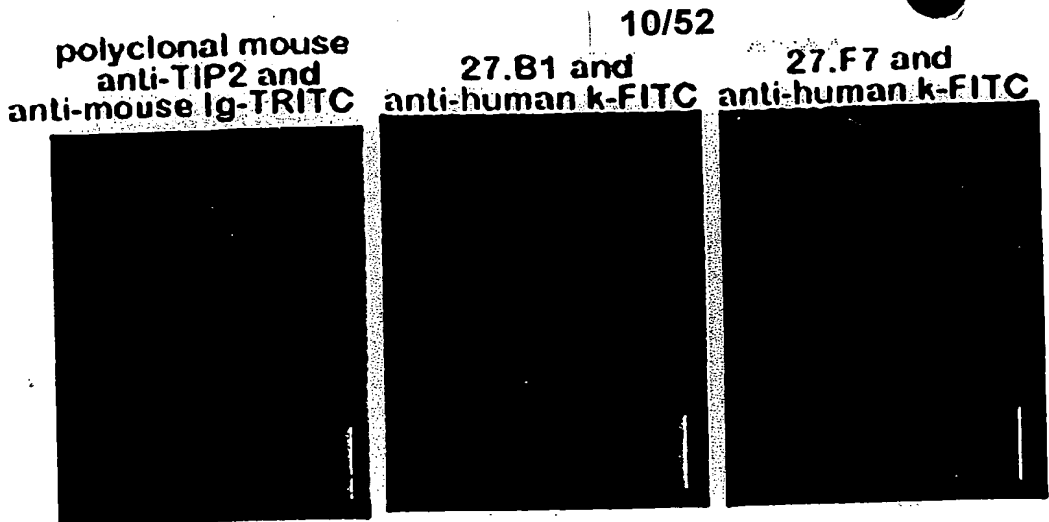
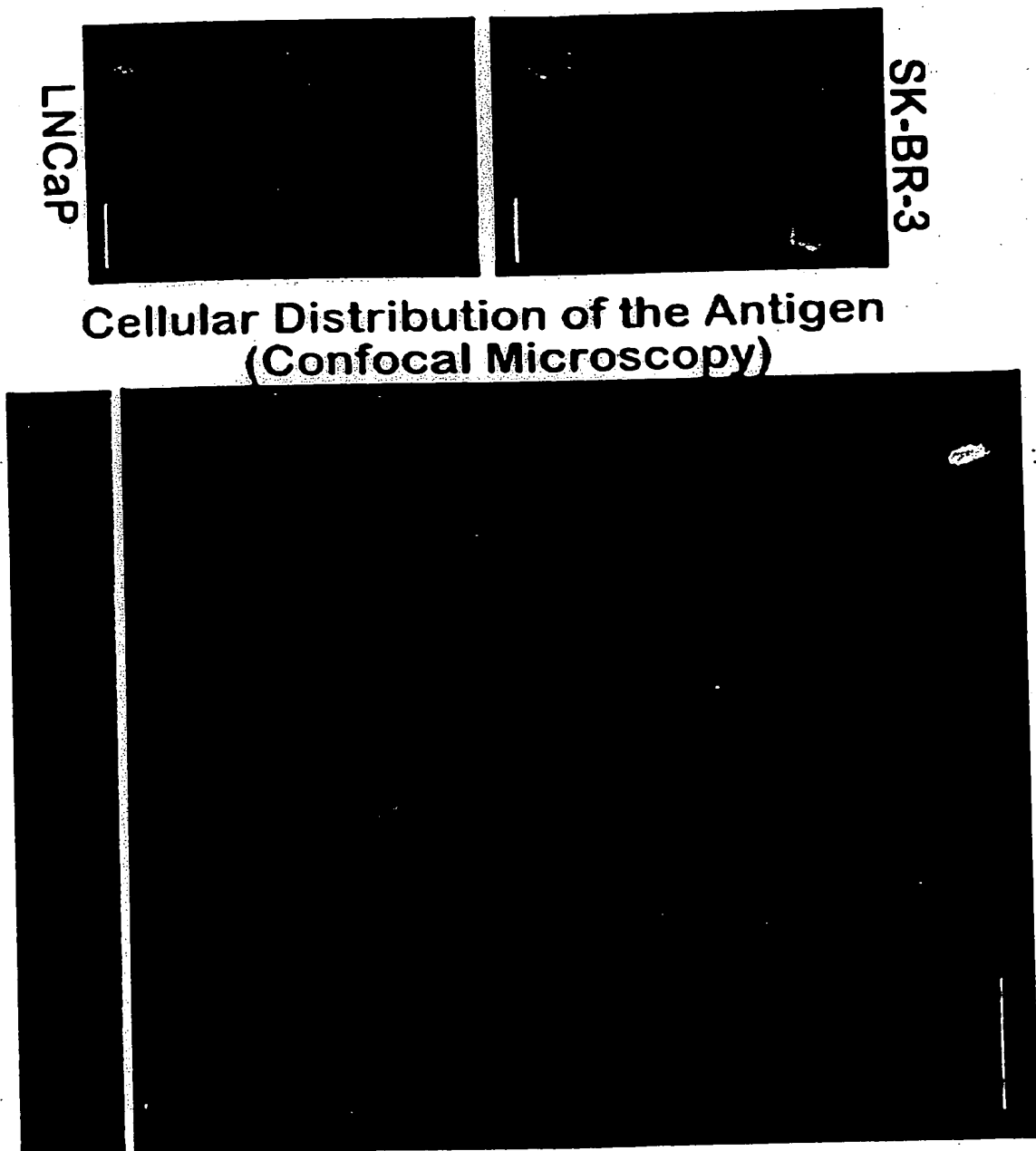


FIG. 9

Detection of TIP2  
in MCF-7 Cells  
using Antibodies



Indirect Immunostaining of Cancer Cells with 27.F7

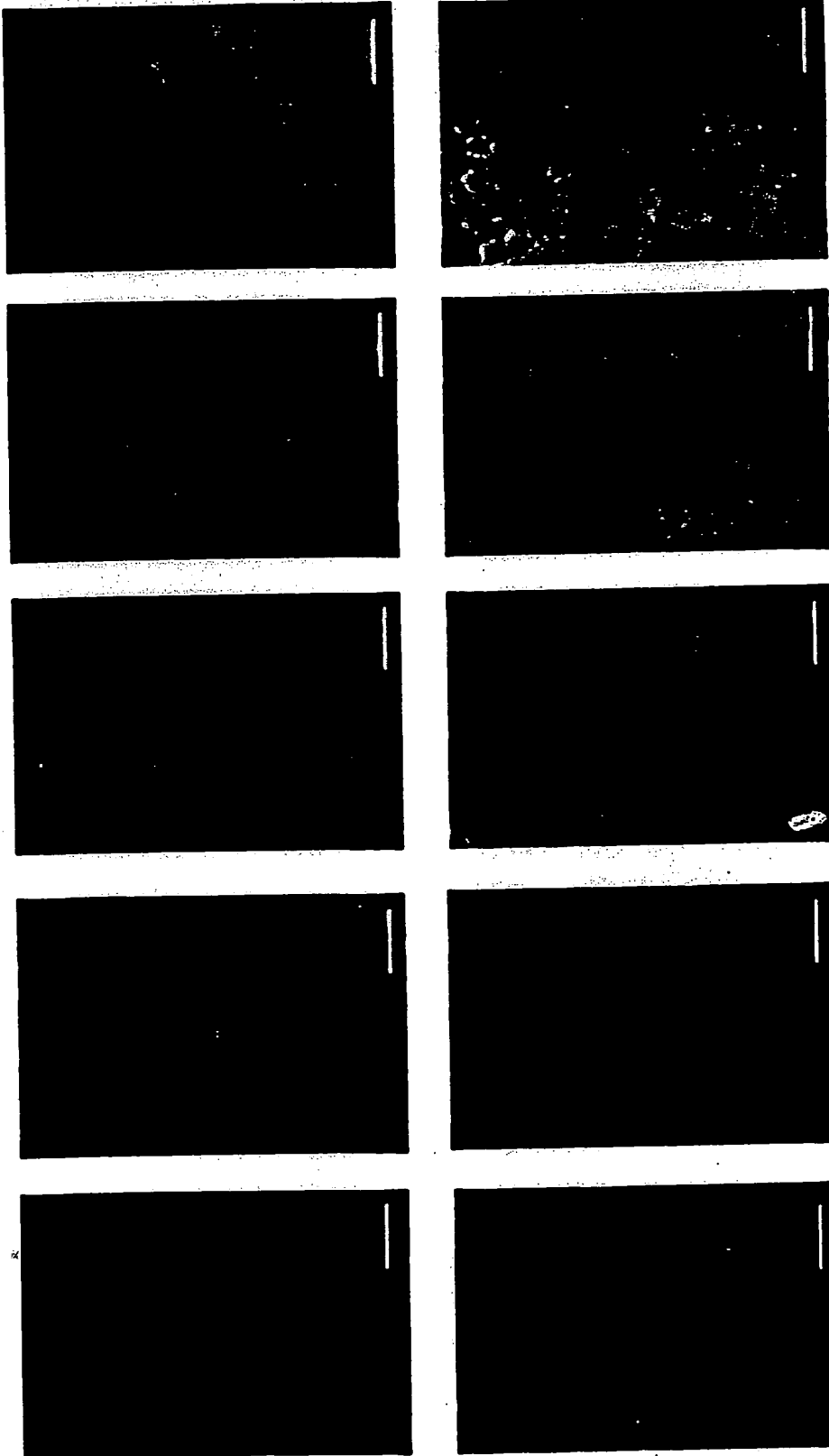


Size bars represent 20  $\mu$ m

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Normal Breast Tissue

Invasive Ductal Cancer



Indirect Immunostaining with 27.B1

FIG. 10

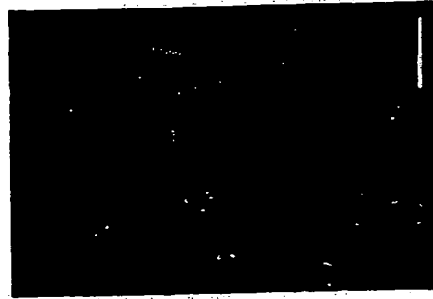
Size bars represent 20  $\mu$ m

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benign prostate hyperplasia

prostate cancer



Indirect Immunostaining with 27.B1

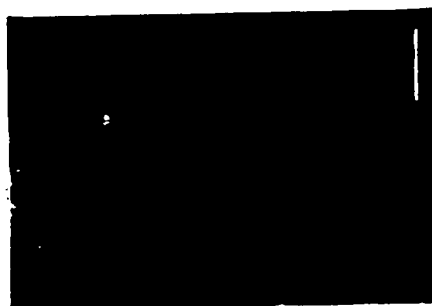
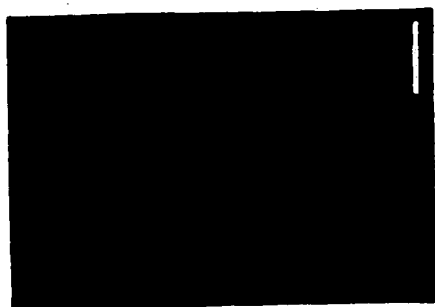
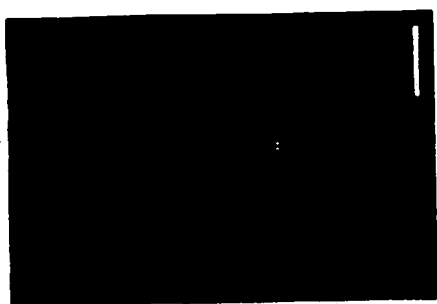
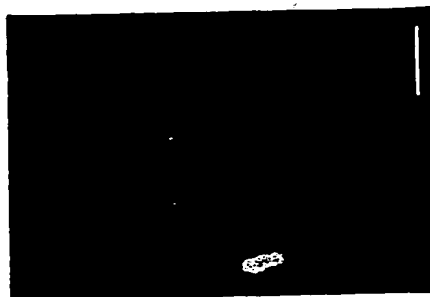
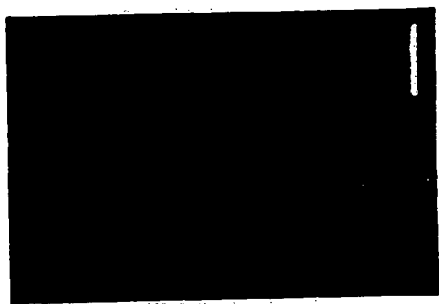
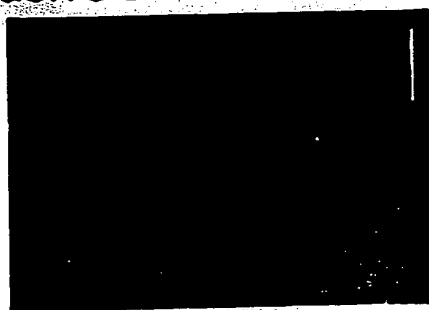
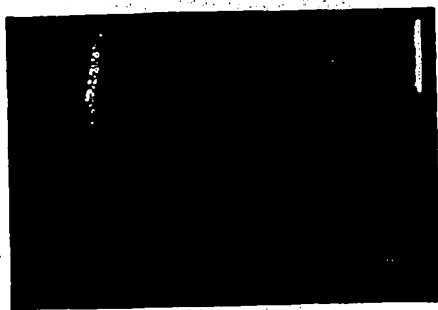
FIG. 11

Size bars represent 20  $\mu$ m  
09664958-091800

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Normal Breast

Invasive Ductal Carcinoma



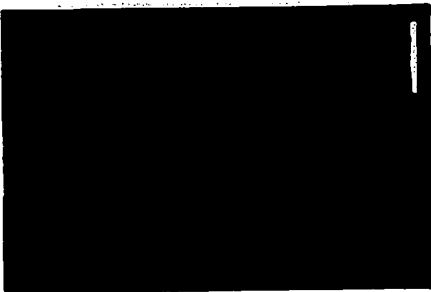
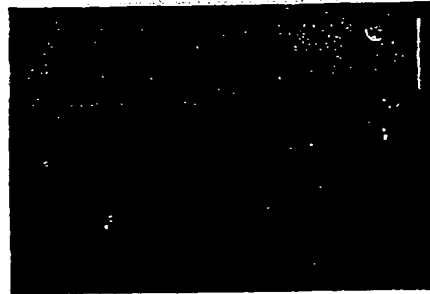
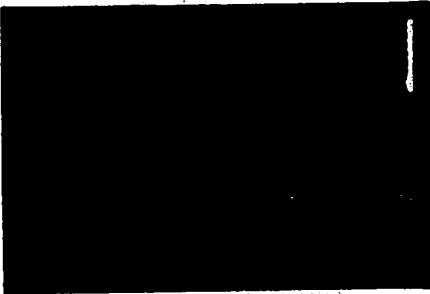
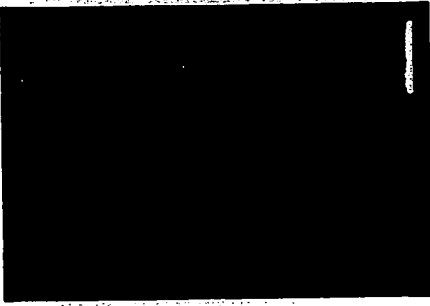
Indirect Immunostaining with 27.F7

FIG. 12

Size bars represent 20  $\mu$ m

benign prostate hyperplasia

prostate cancer



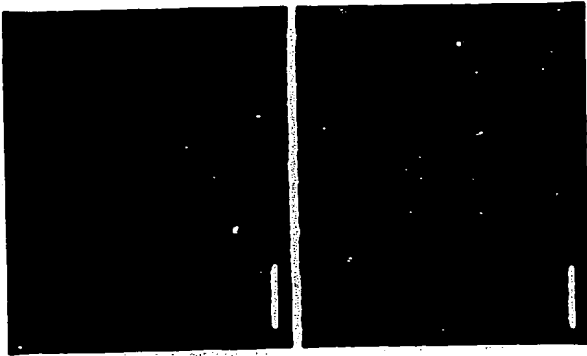
Indirect Immunostaining with 27.F7

FIG. 13

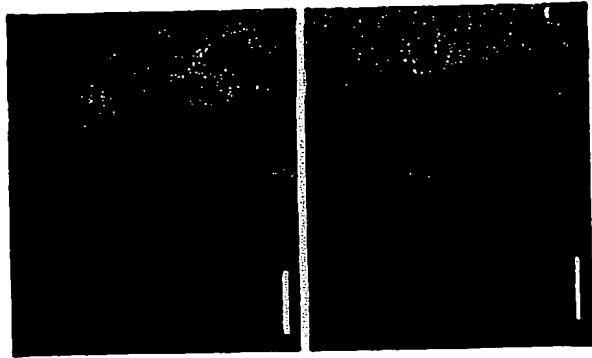
Size bars represent 20  $\mu$ m

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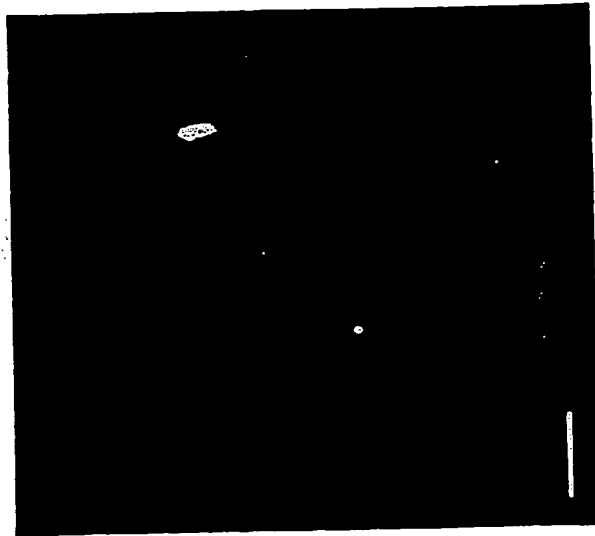
Antibody: 27.F7



Antibody: 27.B1



Distribution of the Antigen (Confocal Microscopy)



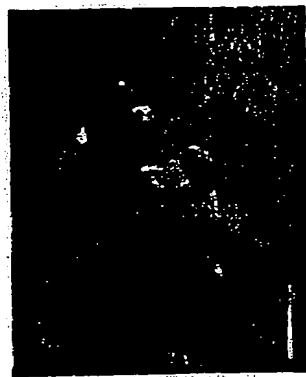
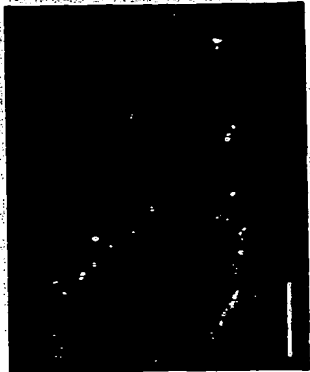
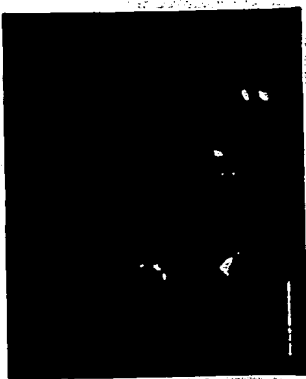
Immunostaining of Breast Cancer Metastases  
in Regional Lymph Nodes

FIG. 14

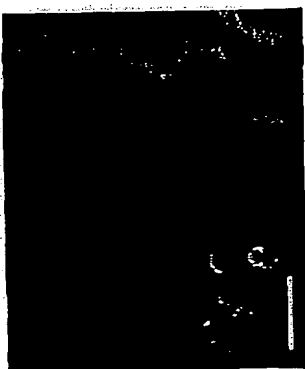
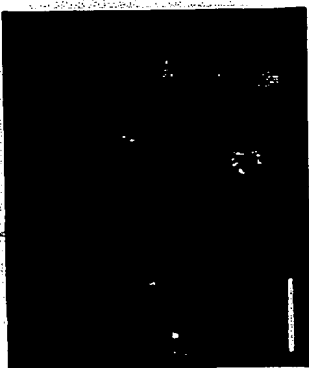
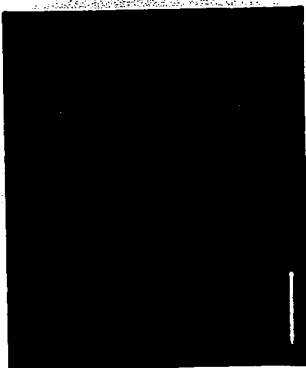
Size bars represent 20  $\mu$ m

Size bars represent 20  $\mu$ m

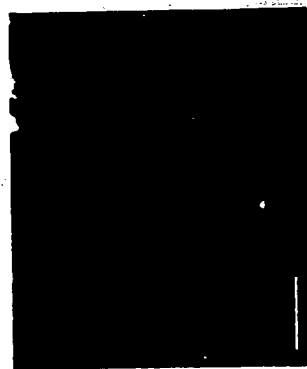
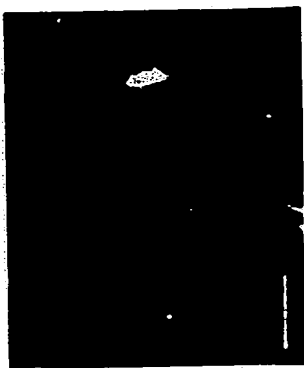
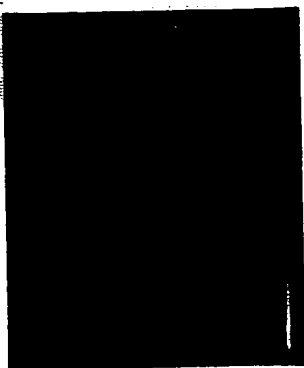
frozen



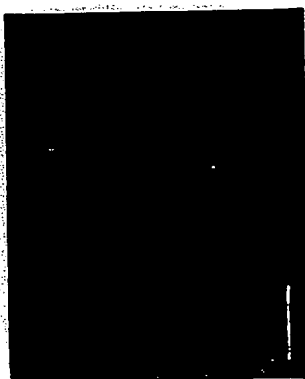
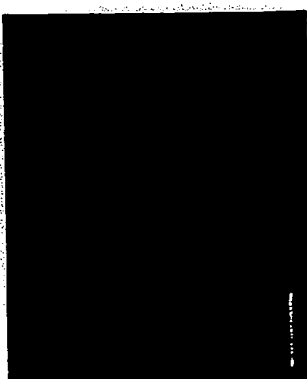
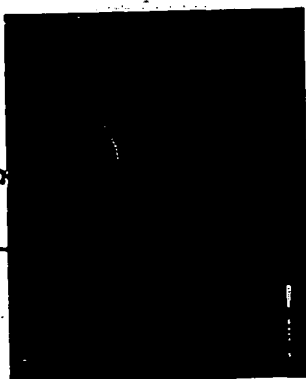
fixed



frozen



frozen



fixed

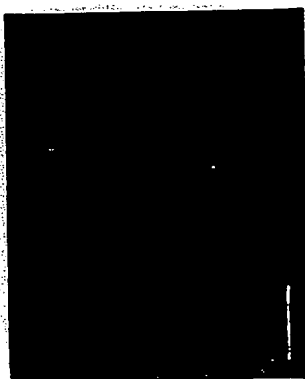
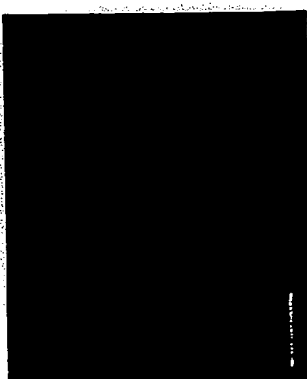
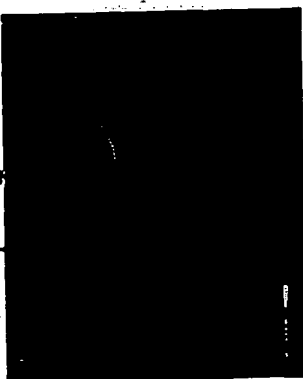


FIG. 15

Indirect Immunostaining of Invasive Ductal Cancer with  
27B1  
27F7



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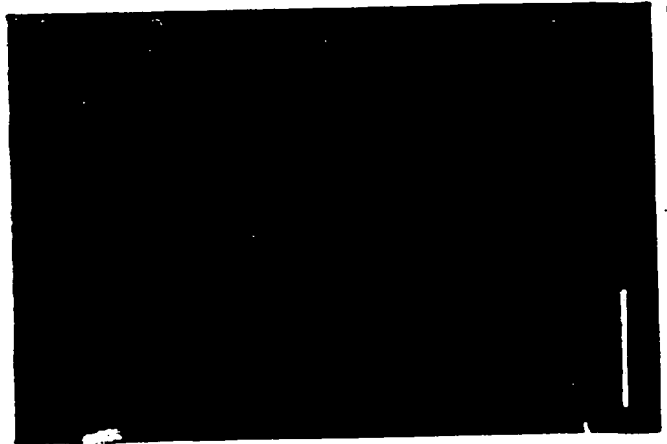
27.F7



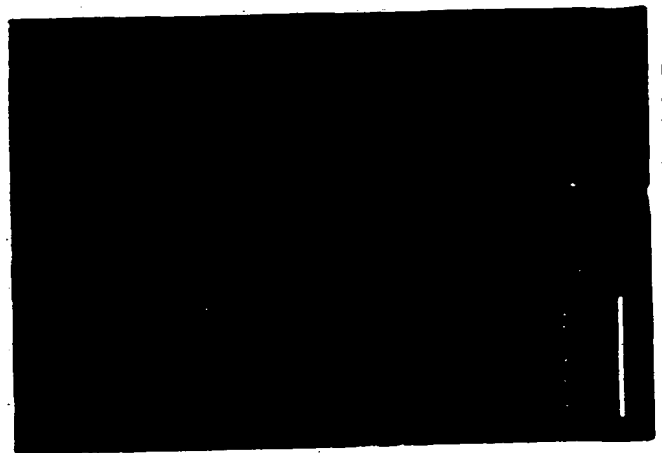
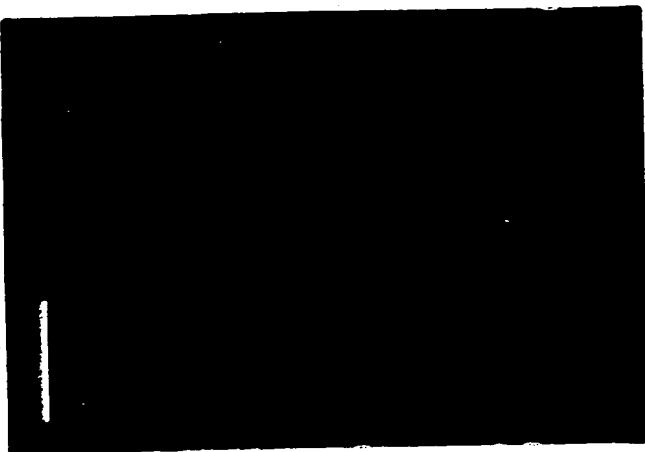
27.B1



IMAGI INUTAUUICIAI VAI UINIVINIA



UUNIVINIA



Size bars represent 20  $\mu$ m

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Indirect Immunostaining with 27.B1

FIG. 17

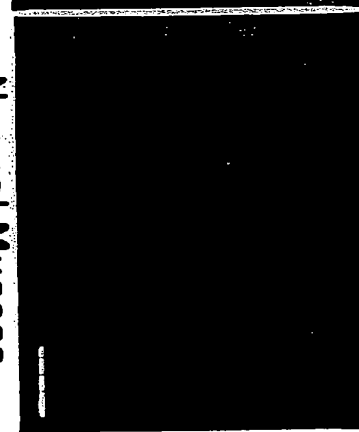
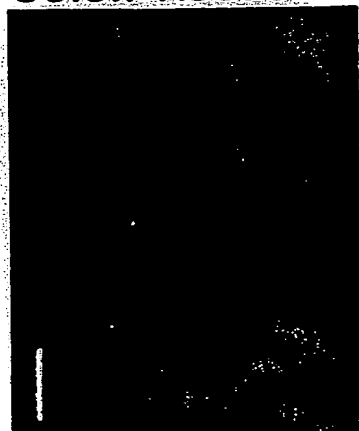
Breast Cancer Tissue



Invasive Ductal Cancer

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Colon Tissue



Colon Cancer

Normal Mucosa

Lung Tissue



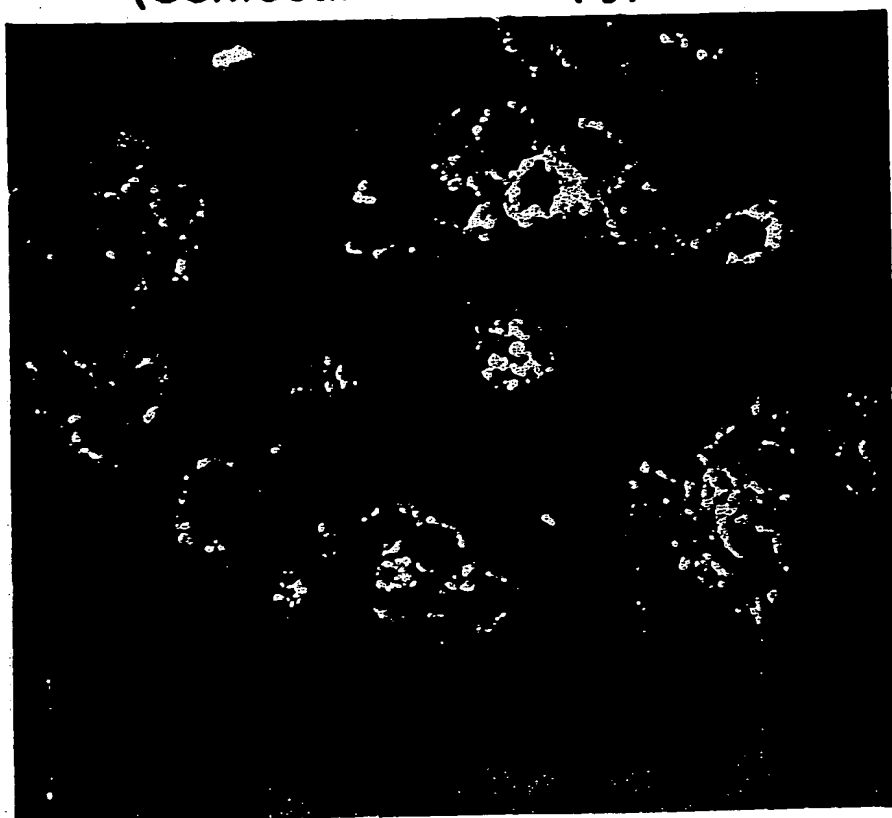
Adenocarcinoma

Non-Cancerous Alveoli

Normal Liver Tissue

Normal Blood

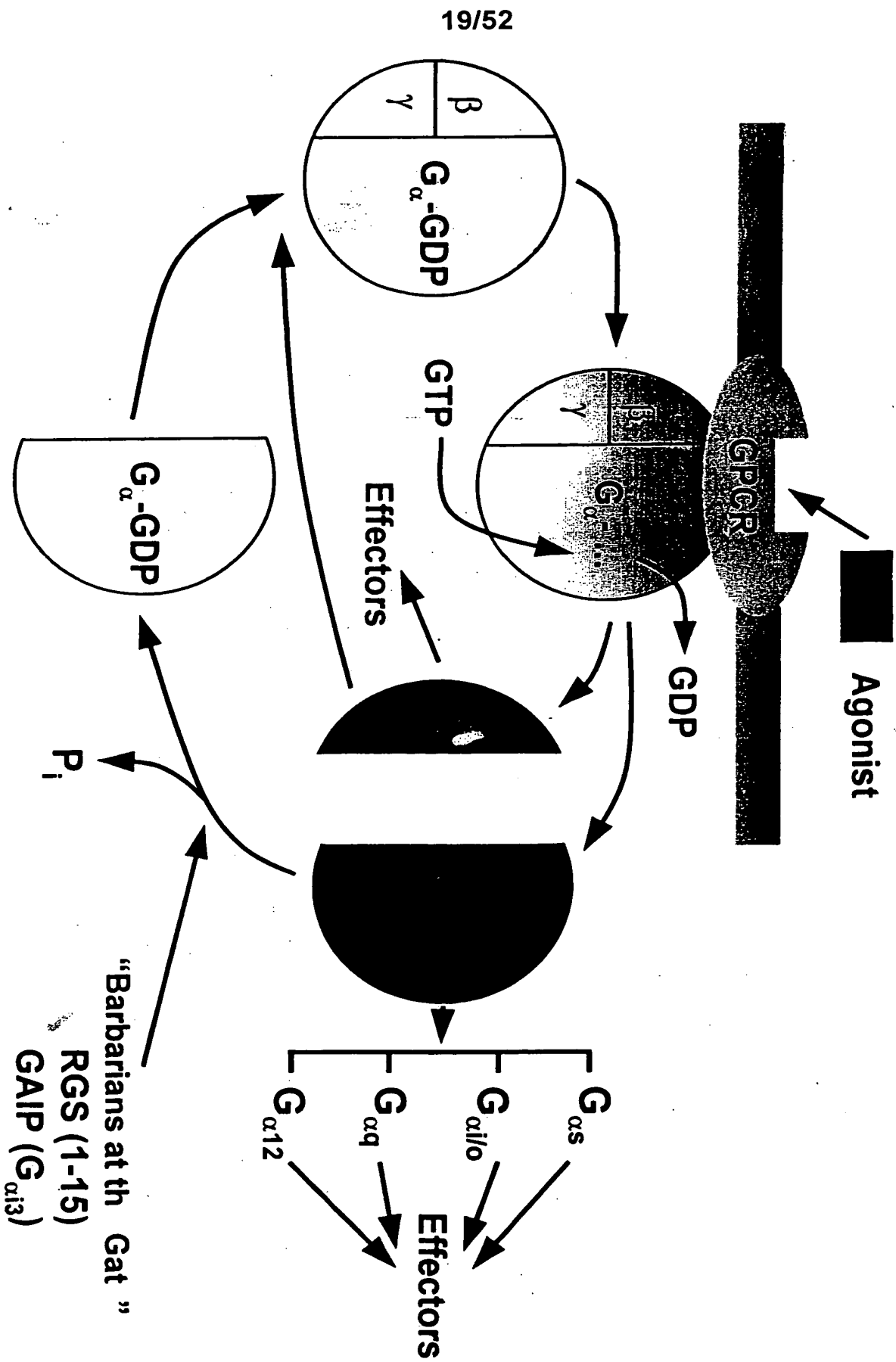
Distribution of the Antigen  
(Confocal Microscopy)



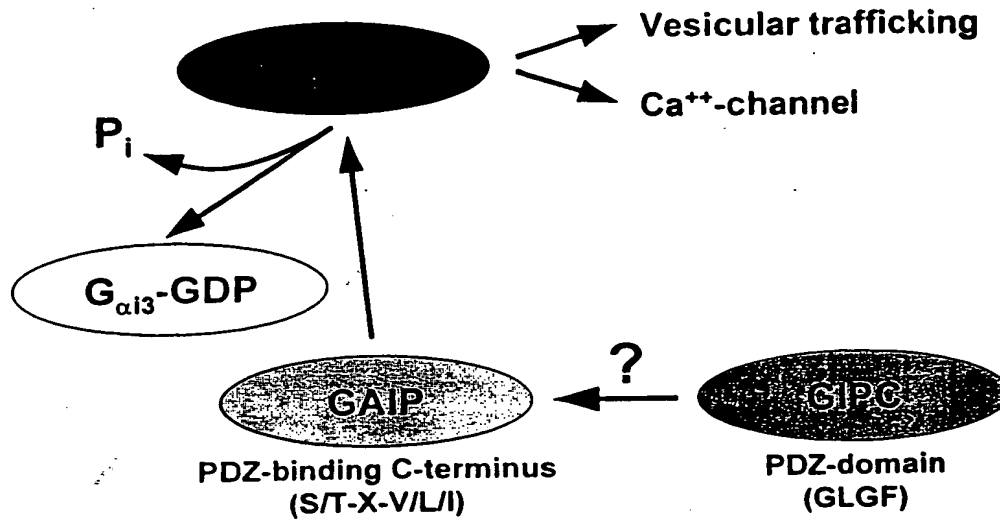
Size bars represent 20  $\mu$ m

FIG. 18

# Regulation of G-protein Signaling System



**FIG. 19 GIPC Proteins (GAIP Interacting Protein, C-terminus) - Regulators of Regulators?**



#### GIPC Family Proteins

- TAX interacting protein 2 (TIP-2)
- Neurophilin binding protein (NIP)
- M-Semaphorin F cytoplasmic domain associated protein (SEMCAPI-1)

#### Other PDZ-"binders"

- NMDA
- TAX oncoprotein
- HPV E6
- AdD9 E4
- glycoporphin C
- FAS
- APC
- LET-23
- CXCR2 (IL-8 RB)
- CXCR5 (coreceptor HTLV-1/HIV)

#### Other PDZ-"containers"

- PSD-95
- DlgA/DLG
- ZO-1
- p55
- LIN7
- PTPL1/FAP1
- RGS12
- PDZ-73 (NYCO38)

FIG. 20

# PRINCIPLE OF SEROLOGICAL RECOMBINANT EXPRESSION CLONING (SEREX) TECHNOLOGY FOR IDENTIFICATION OF TUMOR ASSOCIATED ANTIGENS

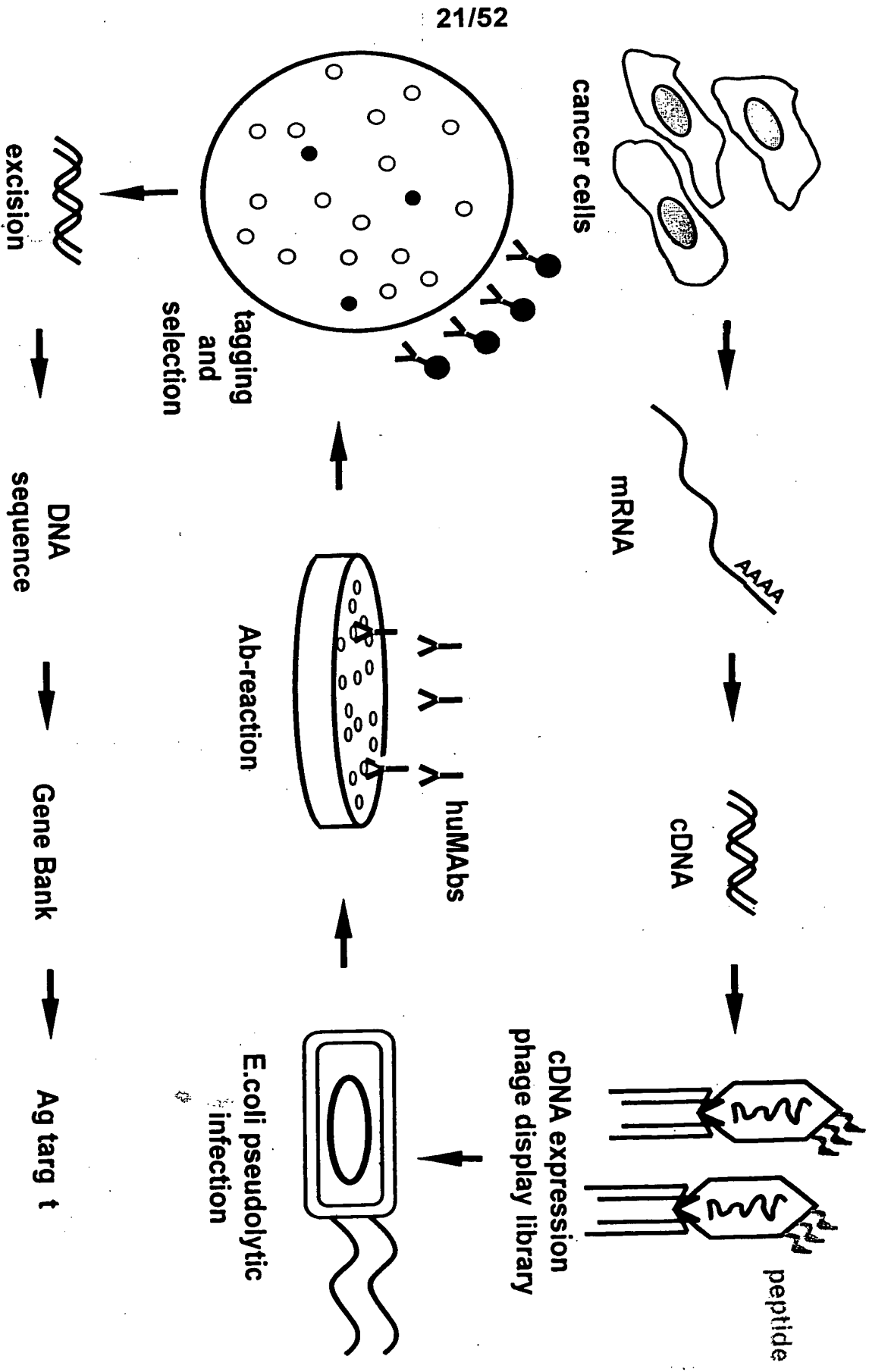
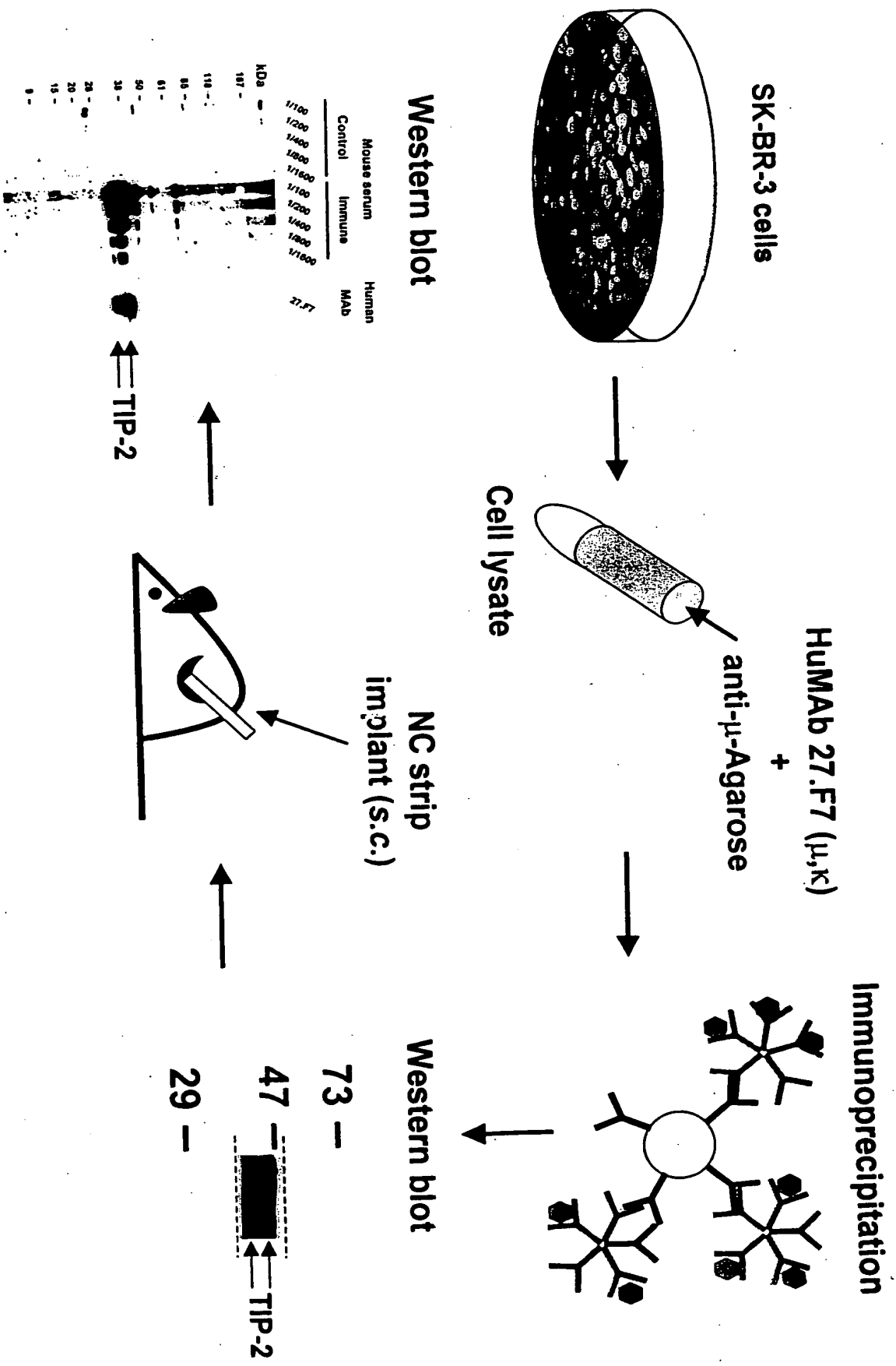


FIG. 21

# DEVELOPMENT OF MOUSE anti-TIP-2 ANTIBODIES USING HUMAN anti-TIP-2 ANTIBODY BOTH AS A CAPTURE AND A TAG

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FIG. 22

SERUM IMMUNOREACTIVITY IN MOUSE IMMUNIZED WITH BREAST CANCER -  
ASSOCIATED ANTIGEN TIP-2

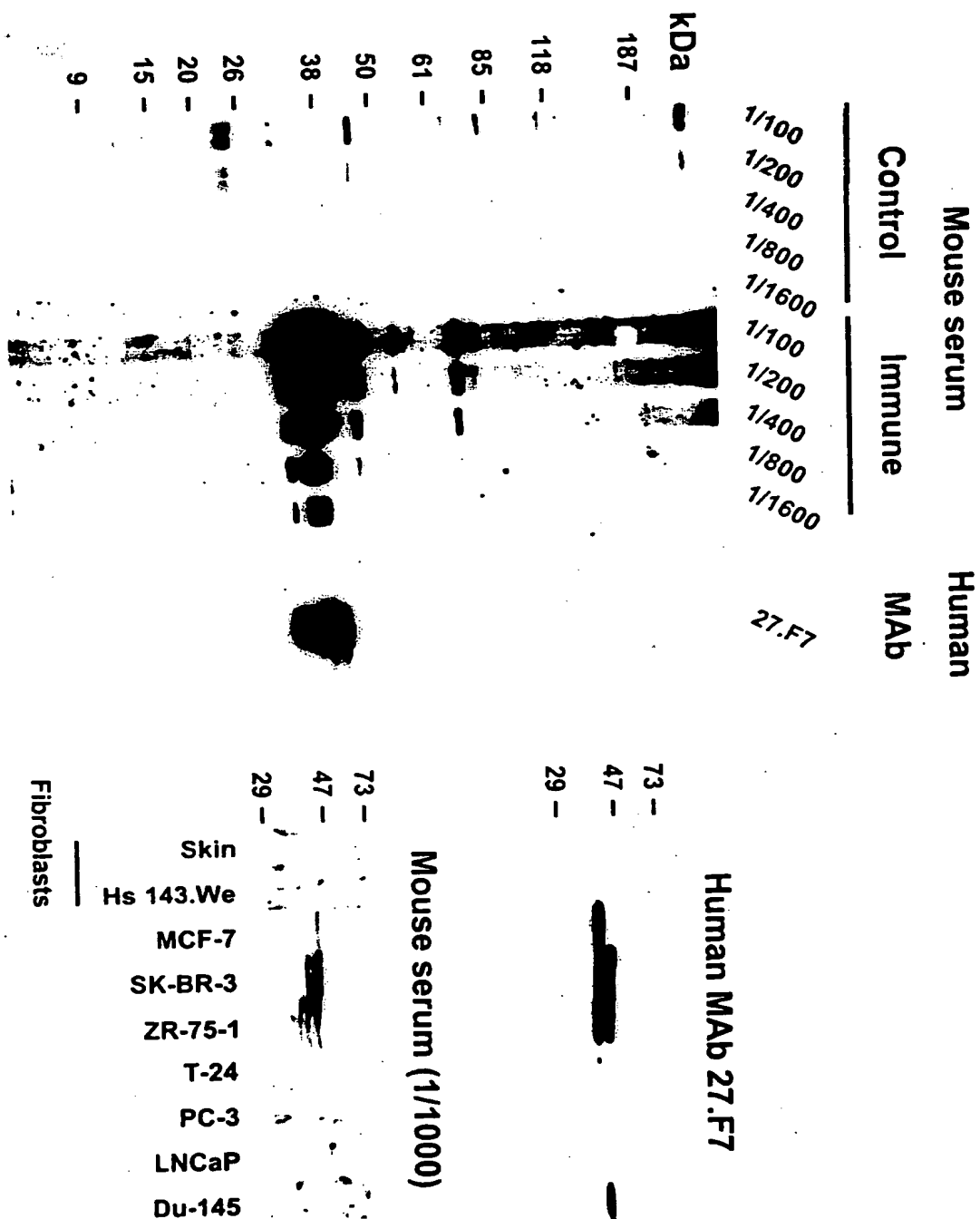


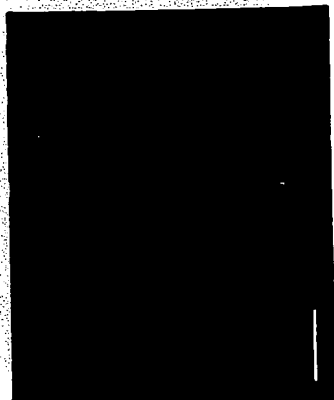
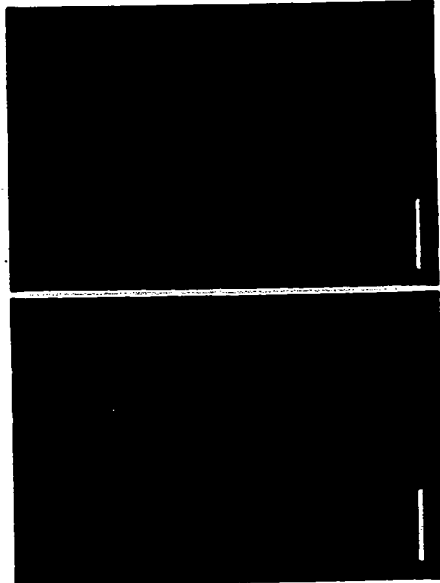
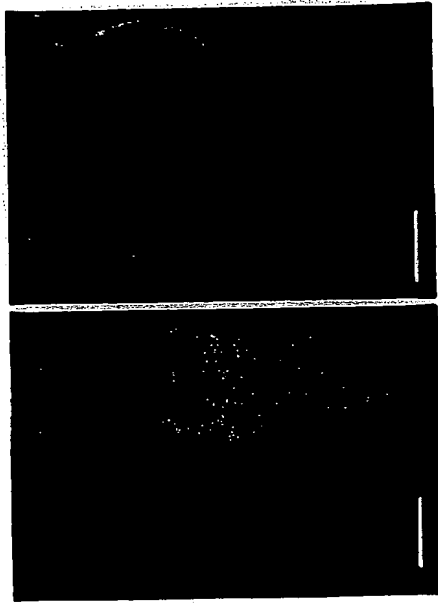
FIG. 23

Invasive Ductal Cancer Tissue Stained Indirectly with:

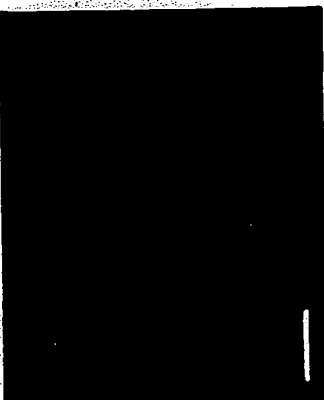
27.F7

polyclonal mouse anti-TIP2

Controls

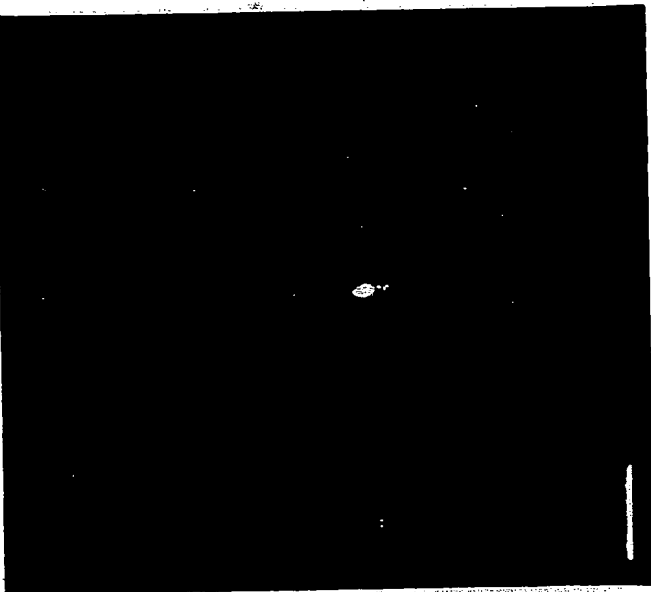
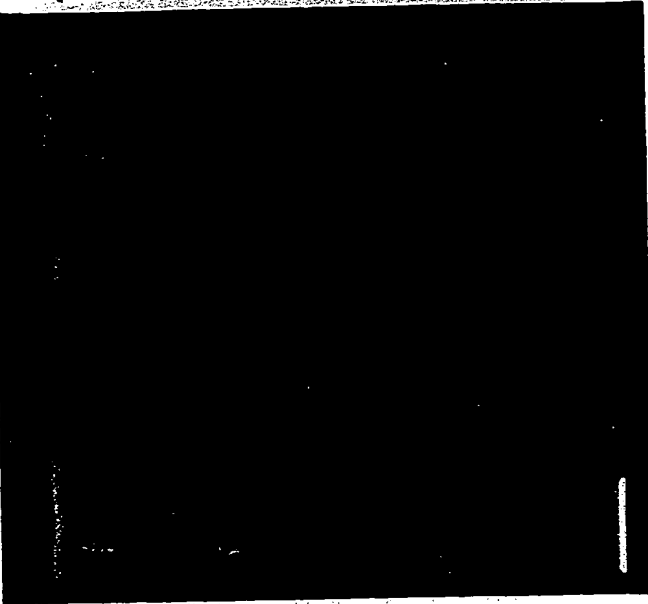


Second Antibody Control



Control Mouse Serum and  
Second Antibody Control

Distribution of the Antigen  
(Confocal Microscopy)



Normal Breast Tissue  
Indirectly stained with  
mouse anti-TIP2

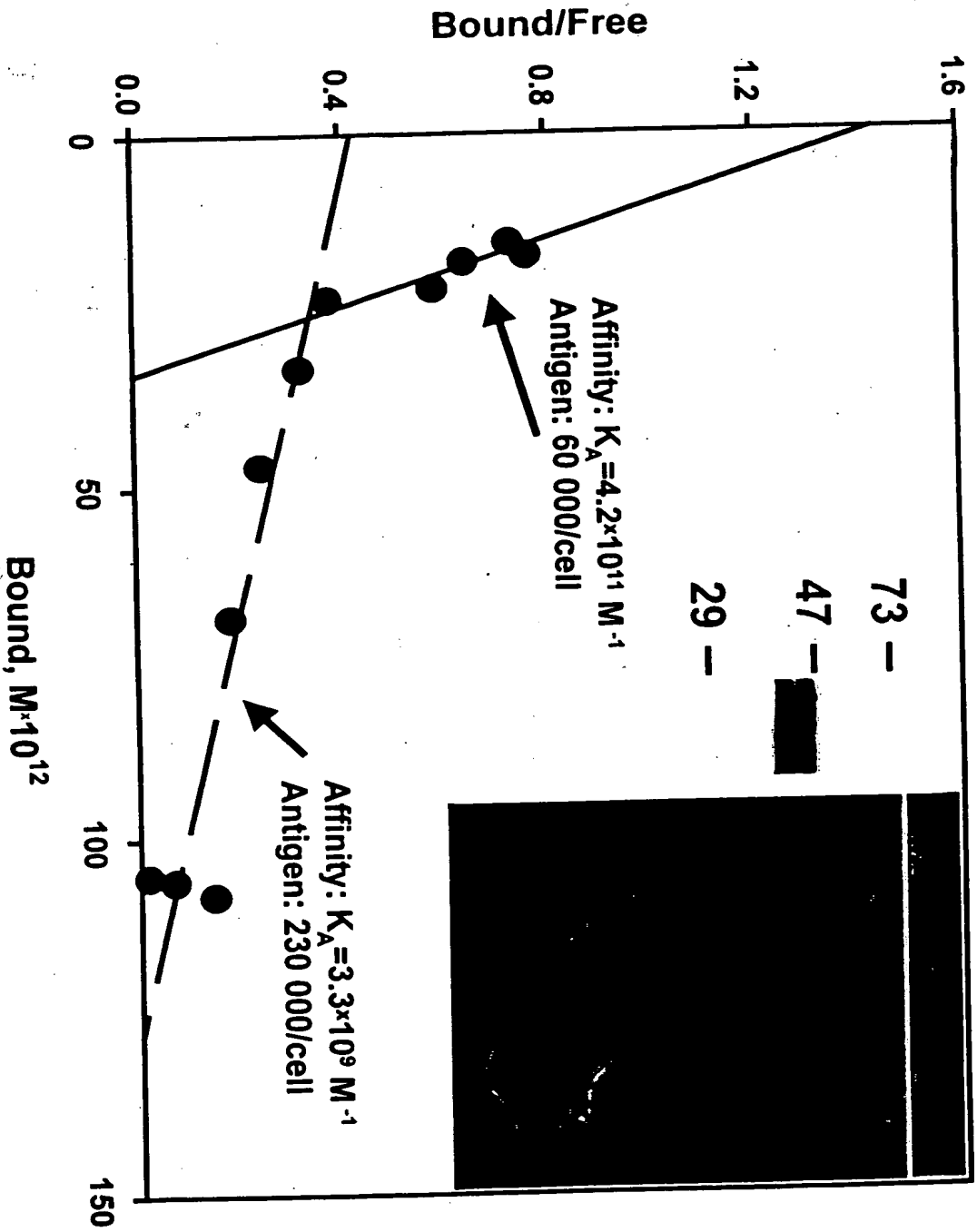
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Size bars represent 20  $\mu$ m



FIG. 24

Analysis for Human anti-TIP-2 Antibody 27.F7 ( $\mu$ ,  $\kappa$ ) on SK-BR-3 Cells



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**FIG. 25 Expression of TIP-2 in Normal and Cancer Breast Tissue Lysates**

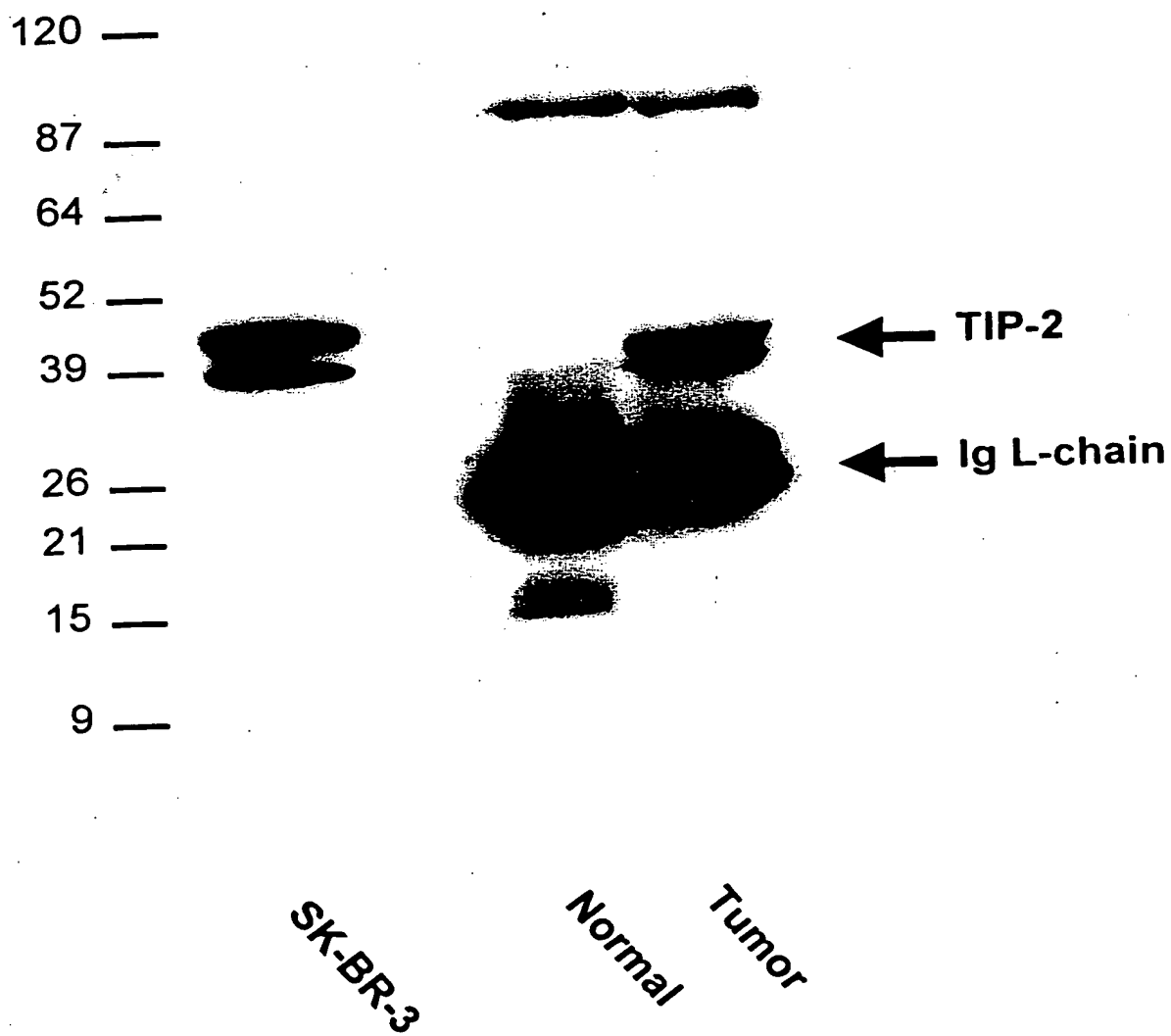
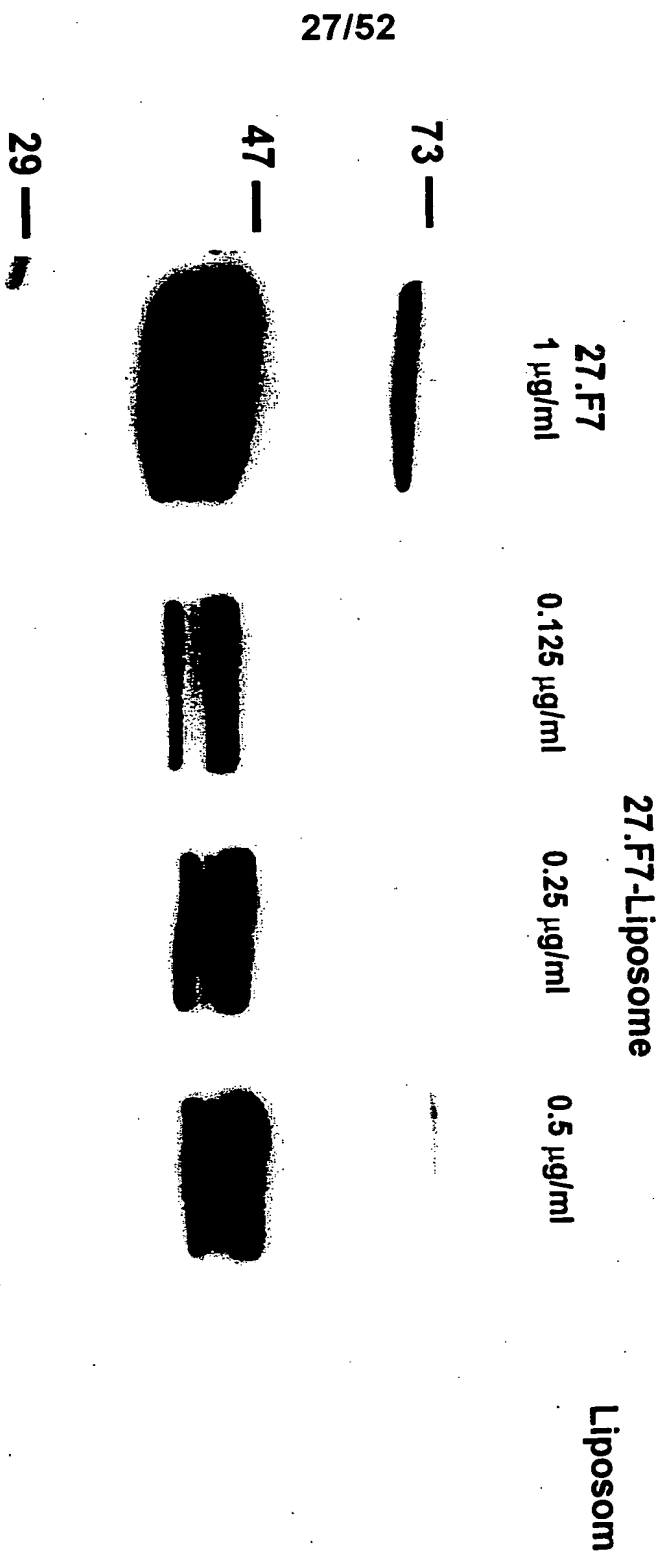


FIG. 26

# Coupling of anti-TIP-2 Antibody 27.F7 ( $\mu$ , $\kappa$ ) to Liposomes

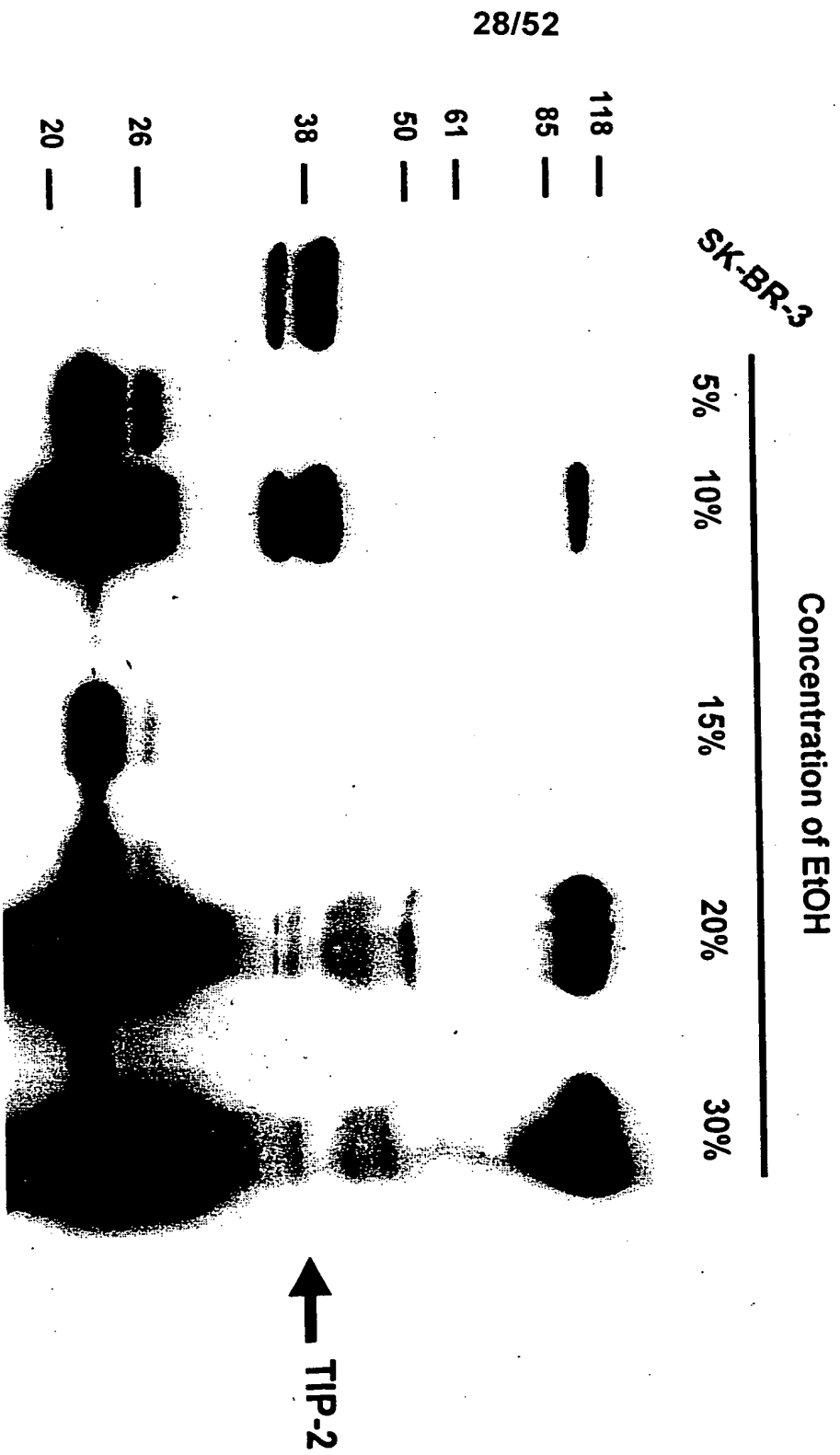


Western blot of SK-BR-3 cell lysate

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FIG. 27

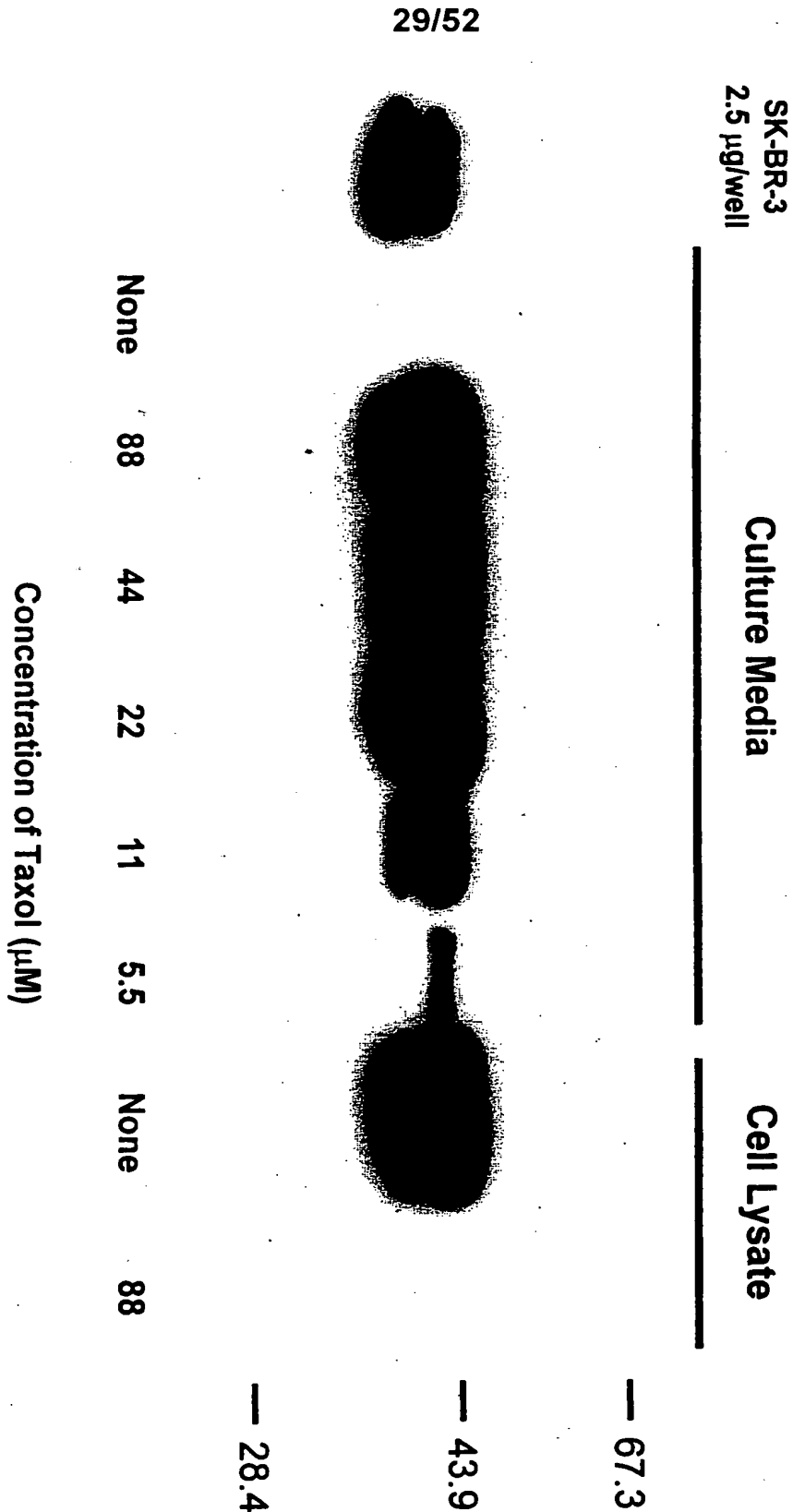
Alcohol Fractionation of Human Serum Spiked  
with SK-BR-3 Lysates (TIP-2 Containing)



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FIG. 28

# Release of TIP-2 into Culture Media from SK-BR-3 Cells Treated by Taxol



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FIG. 29

# Amino Acid Sequence of GLUT1CBP/GIPC Protein

10	20	30	40	50	60
MEPLGLRRKK	APPLVENEFA	EPGRGLGVG	EPGLGGGS	GGPQGLPPP	PPALRPLVF
70	80	90	100	110	120
HTQLAHGSPT	GRIEGFTNVK	EIYKIAEAF	RLPTAEVMFC	TLNTHKVDMD	<u>KLGGQIGLE</u>
130	140	150	160	170	180
DFIEAHVKGQ	RKEVEVERKSE	DALGLTTDN	GAGYAFIKRI	KEGSVIDHIH	LISVGMIEA
190	200	210	220	230	240
<u>INGQSLGCR</u>	<u>HYEVARLIKE</u>	LPRGRFTYIK	LTEPRKAFDM	ISQRSAGRP	GSQPQLGTGR
250	260	270	280	290	300
GTIRLRSRGP	ATVEDLPSAF	EKAIEKVD	LLESYMGIRD	TELAATMVEL	GKDKRNPDEL
310	320	330			
AEALDERLGD	FAFPDEFVFD	VWGAIGDAKV	GRY		

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TIP-2 sequence is shown in *italic*

HLA A\*0201 binding peptides (111-119 and 185-194) are shown underlined

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FIG. 30

1 caccgggagc cggaggcagc ggcggcgccg gcggcgccg cggcgccgc ggaagcagatc  
61 ttctgtgtgac cccactctc gctgtcatg ccgtctggac tggggcgccg gaaaaagcgc  
121 cccctctag tggaaaatga ggaaggctgag ccaggccgtg gaaggctggg cgtgggggag  
181 ccagggcctt tgggcggag tgggtcgggg ggccccaaaa tgggcttgcc cccctccc  
241 ccaggccctgc ggcgccgcct tgtgtccac acccagctgg cccatggcag tcccatggc  
301 cgcatacgag ggttcaccaa cgtcaaggag ctgtatggca agattgccga ggccttcgc  
361 ctgccaaactg ccgaagtgat gttttgacc ctgaacaccc acaaatgga catggaacaag  
421 ctccctggggg gccaaatcgg gctggaggac ttcatcttcg cccacgtgaa ggggcaagcgc  
481 aaggaggtgg aggtgttcaa gtcggagagat gcactcgggc tcaaccatcac ggaacaaagg  
541 gctggctacg ccttcacaa gcccatcaag gaaggcagcg tgatcgacaa catccaactc  
601 atcagcgtgg gcgacatgat cgaagccatt aacgggcaga gccctgtggg ctgccggcac  
661 tacgaagtgg cccgctgct caaggaaactg ccccgaggcc glacctcac gctgaagctc

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FIG. 31

Protein Antigens Identified by Natural Human Monoclonal Antibodies Developed from Breast and Prostate Cancer Patients' B-Cells

Antibody	Antigen Name	Sequence	Molecular Weight (Calculated)	HLA A*0201-Specific MHC Binding Peptides	mRNA Expression in Tissues	Functions
13.42 $\mu$ κ	Human mRNA for KIAA0338 gene, partial cds	See Fig. 32	103568 (~40KD by WB)	NLEKDYFGL (184-193) VLFDLVCEHL (174-183) KLQHPDMLV (903-911)	Brain	Unknown
13.2C1 $\mu$ κ	Human non-muscle alpha-actinin mRNA, complete cds - the second non muscle alpha-actinin isoform designated ACTN4 (actinin-4)	See Fig. 33	105217	KMLDAEDIV (238-246) KMTLGMIVTI (139-148) FMPSGKMOV (374-382) KLASDLLEWI (302-311) GLVTFQAFI (825-833) CQLEINFNSV (353-362)	Adipose, Adrenal gland, Aorta, Brain, Breast, CNS, Colon, Ear, Esophagus, Foreskin, Germ Cell, Heart, Kidney, Liver, Lung, Muscle, Ovary, Pancreas, Parathyroid, Placenta, Prostate, Small intestine, Stomach, Testis, Thyroid, Tonsil, Uterus, Whole embryo, breast, colon, genitourinary tract, head_neck, lung, cell line, ovary, stomach	Actin-binding protein important in organization of cytoskeleton and in cell adhesion. "An amino-terminal fragment of alpha-actinin can promote monocyte/macrophage maturation" [Exp. Hematol. 1999, 27(2):345-52].
13.2C1 $\mu$ κ	Homo sapiens actinin, alpha 4 (ACTN4) mRNA	See Fig. 34	102260	KMLDAEDIV (212-220) KMTLGMIVTI (113-122) FMPSGKMOV (345-353) KLASDLLEWI (273-282) GLVTFQAFI (797-805)	Adipose, Adrenal gland, Aorta, Brain, Breast, CNS, Colon, Ear, Esophagus, Foreskin, Germ Cell, Heart, Kidney, Liver, Lung, Muscle, Ovary, Pancreas, Parathyroid, Placenta, Prostate, Small intestine, Stomach, Testis, Thyroid, Tonsil, Uterus, Whole embryo, breast, colon, genitourinary tract, head_neck,	Actin-binding protein important in organization of cytoskeleton and in cell adhesion. "The cytoplasmic localization of actinin-4 was closely associated with an infiltrative histological phenotype and correlated significantly



					lung, cell line, ovary, stomach	with a poorer prognosis in 61 cases of breast cancer" [J. Cell. Biol. 1998, 140(6):1383-93]. Alpha-actinin-1 and 4 associate with PDZ domain of CLP-36 PDZ-LIM protein (also called hCLIM1 - high expression in epithelial cells) in actin stress fibers [JBC 2000, 275(15):11100-11105].
22.8D11 $\mu\lambda$	Human clathrin coat assembly protein 50 (AP50) m RNA	See Fig. 35	49662	WLAAVTKQNV (64-73) ILPFRVPLV (284-293) SLLAQKIEV (314-322) KLNYSDHDV (410-418)	infant brain, brain, placenta, breast, ovary (tumor), fetal heart, fetal lung, multiple sclerosis lesions, pineal gland, lymph node	Component of the adaptor complexes which link clathrin to receptors in coated vesicles clathrin-associated protein complexes are believed to interact with the cytoplasmic tails of membrane proteins, leading to their selection and concentration. AP50 is a subunit of the plasma membrane adaptor.
27.B1 $\mu\kappa$ 27.F7 $\mu\kappa$	Homo sapiens GLUT1 C-terminal binding protein (GLUT1CBP) mRNA [GIPC/TIP-2]	See Fig. 36	36047	KLGGQIGL (111-119) SLGGRHYEV (185-194)	Adipose, Aorta, Blood, Bone, Brain, Breast, CNS, Colon, Germ Cell, Heart, Kidney, Lung, Ovary, Pancreas, Placenta, Pooled, Stomach, Testis, Thymus, Uterus, Whole embryo, brain, breast, colon, connective tissue, lung, muscle	Binds via a PDZ domain to C terminus of GLUT1 and interact with cytoskeletal proteins
33.2H6 $\mu\lambda$	Homo sapiens gpl30 associated protein GAM mRNA	See Fig. 37	21835	YLSQEHQQQV (94-103)	placenta, breast, infant brain, uterus (pregnant), B-Cell, ovary (tumor), fetal heart, fetal liver/spleen, fetal lung, T cells (Jurkat cell line)	Has a possible role in the negative regulation of proteins containing WD-40 repeats. May be required for the initiation and maintenance of the differentiated state.

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33.2H6 $\mu\lambda$	Homo sapiens amino-terminal enhancer of split (AES) mRNA	See Fig. 38	21966	YLSQEHQQQV (95-104)	Adrenal gland, Aorta, Blood, Bone, Brain, Breast, CNS, Colon, Esophagus, Eye, Foreskin, Germ Cell, Head and neck, Heart, Kidney, Lung, Lymph, Muscle, Nose, Ovary, Pancreas, Parathyroid, Placenta, Pooled, Prostate, Spleen, Stomach, Synovial membrane, Testis, Thymus, Thyroid, Tonsil, Uterus, Whole embryo, brain, colon, head_neck, kidney, lung, ovary, pnet	Amino-terminal enhancer of split is similar to the Drosophila enhancer of split groucho protein. The function of AES has not been determined but it has been proposed as a candidate tumor human cancer antigen.
33.2H6 $\mu\lambda$	Antiquitin 1 (antiquitin=26g turgor protein homolog), mRNA	See Fig. 39	55357	KVMDRPGNYV (372-381) ALIEQWNPV (149-157) IITAFNFPV (162-170)	fetal heart, infant brain, placenta, NT2 neuronal precursor, liver, Hel a (cell line), ovary, liver (HepG2 cell line), ovary (tumor), multiple sclerosis lesions	Unknown (30% identity to various eukaryotic and prokaryotic aldehyde dehydrogenases). Antiquitin has homology to a previously described protein from the green garden pea, the 26g pea turgor protein. Four human antiquitin-like sequences, possibly pseudogenes, have also been identified.
39.A7 $\mu\lambda$	ARP2/3 protein complex 41 KD subunit (P41-ARC), mRNA	See Fig. 40	40935	FEQENDWVV (125-133)	Hel a (cell line), fibroblast, fetal brain, infant brain, fetal liver/spleen, monocytes (stimulated), fetal heart, uterus (pregnant), olfactory epithelium, breast	Part of a complex implicated in the control of actin polymerization in cells belongs to a complex composed of ARP2, ARP3, P41-ARC, P34-ARC, P21-ARC, P20-ARC and P16-ARC.
50.1B3 $\mu\kappa$	H.sapiens seb4D mRNA H.sapiens seb4B mRNA	See Fig. 41a and 41b	seb4D-24617	for seb4D YLGAKPWCL (100-108) CLQTGFAIGV (107-116)	thymus, Blood, Brain, Breast, Colon, Germ Cell, Heart, Kidney, Lung, Lymph, Ovary, Parathyroid, Pooled, Prostate, Testis, Thymus, Tonsil, Uterus, brain, colon, lung, muscle, ovary,	Unknown

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			seb4B- 25218	for seb4B YLGAKPWCL (101-109) CLQTGFAIGV (108-117)	stomach, thymus, pooled, whole blood	
59.3G7 μλ	Homo sapiens lamin A/C (LMNA) mRNA	See Fig. 42	65133	KLLEGEERL (378-387) KLVRSVTVV (542-550) RLADALQEL (240-248)	Adipose, Adrenal gland, Bone, Brain, Breast, Colon, Esophagus, Foreskin, Germ Cell, Heart, Kidney, Larynx, Liver, Lung, Lymph, Muscle, Ovary, Pancreas, Parathyroid, Placenta, Pooled, Prostate, Spleen, Stomach, Synovial membrane, Testis, Thymus, Thyroid, Uterus, Whole embryo, brain, breast, colon, denis_drash, head_neck, lung, cell line, ovary, stomach	Intermediate filament proteins

## FIG. 32

Human mRNA for KIAA0338 gene, partial cds

ORIGIN

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1 catcagcggg cgggggtgtc gccgaacagg ctgctccgca gagcccgcg cgaccccgcg
61 ccgccccgcc ccgcggcctg cctgccagag gagccgaggg ggccgcccct cgcccaacct
121 gcccgcacatg gggaaccccg ggcccaggcg tgctggtcac catgacaaca gagacaggcc
181 ccgactctga ggtgaagaaa gctcaggagg aggccccgca gcagcccagag gctgctgccg
241 ctgtgaccac ccctgtgacc cctgcaggcc acggccaccc agaggccaac tccaatgaga
301 agcatccatc ccagcaggac acgcggcctg ctgaacagag cctagacatg gaggagaagg
361 actacagtga ggccgatggc ctttcggaga ggaccacgcc cagcaaggcc cagaaatcgc
421 cccagaagat tgccaagaaa tacaagagtg ccatctgccg ggtcactctg cttgatgcct
481 cggagtatga gtgtgagggtg gagaaacatg gccggggcca ggtgctgttt gacctggtct
541 gtgaacacct caacctccta gagaaggact acttcggcct gaccttctgt gatgctgaca
601 gccagaagaa ctggctggac ccctccaagg agatcaagaa gcagatccgg agtagcccct
661 ggaattttgc cttcacagtc aagttctacc cgcctgatcc tgcccagctg acagaagaca
721 tcacaagata ctacctgtgc ctgcagctgc gggcagacat catcacgggc cggctgccat
781 gctcctttgt cacgcatgcc ctactgggct cctacgctgt gcaggctgag ctgggtgact
841 atgatgctga ggagcatgtg ggcaactatg tcagcgagct cegcttcgcc cctaaccaga
901 cccgggagct ggaggagagg atcatggagc tgcataagac atataggggg atgaccccgg
961 gagaagcaga aatccacttc ttagagaatg ccaagaagct ttccatgtac ggagtagacc
1021 tgcaccatgc caaggactct gagggcatcg acatcatgtt aggcgtttgt gccaatggcc
1081 tgctcatcta ccgggaccgg ctgagaatca accgctttgc ctggcccaag atcctcaaga
1141 tctcctacaa gaggagtaac ttctatatca agatccggcc tggggagtat gagcaatttg
1201 agagcacaat tggctttaag ctcccaaacc accggtcagc caagagactg tggaggtctt
1261 gcatcgagca tcatacatte ttccggctgg tgtcccctga gccccaccc aagggttcc
1321 tggatgatgg ctccaagttc cgtacagtg ggaggacca ggcacagact cgccaggcca
1381 gcgcctcat tgaccggcct gcacccttct ttgagcgctt ttccagcaaa cggtagacca
1441 tgtcccgcag ccttgatgga gcagagttct cccgcccagc ctcggtcagc gagaacctg
1501 atgcagggcc tgacggtgac aagcgggatg aggatggcga gtctgggggg caacgggtcag
1561 aggctgagga gggagaggtc aggactccaa ccaagatcaa ggagctaaag ccggagcagg
1621 aaaccacgcc gagacacaag caggagttct tagacaagcc agaagatgtc ttgctgaagc
1681 accaggccag catcaatgag ctcaaaagga ccctgaagga gcccaacagc aaactcatcc

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008160 0564950 091800

FIG. 32 (cont.)

1741 accgggatcg agactgggaa cgggagcgca ggctgccctc ctcccccgcc tccccctccc  
 1801 ccaagggcac ccctgagaaa gccaatgaga gagcagggtt gagggagggc tccgaggaga  
 1861 aagtcaaacc accacgtccc cgggccccag agagtacac aggcgatgag gaccaggacc  
 1921 aggagagggg cacggtgttc ctgaaggaca accacctggc cattgagcgc aagtgtctca  
 1981 gcatcacggt cagctctacg tctagcctgg aggctgaggt ggacttcacg gtcattgggt  
 2041 actaccatgg cagcgccttc gaagacttct cccgcagcct gcctgagctc gaccgggaca  
 2101 aaagcgactc ggacactgag ggcctgctgt tctcccggga tctcaacaag ggggccccca  
 2161 gccaggatga tgagtctggg ggcatgagg acagcccgga tcgaggggccc tgctccaccc  
 2221 cggatatgcc ccagtttgag cccgtgaaaa cagaaacat gactgtcagc agtctggcca  
 2281 ttagaaaaga gattgagccg gaggccgtac tgcagaccag agtctccgct atggataaca  
 2341 cccagcaggt tgatgggagt gcctcagttg ggagggagtt catagcaacc actccctcca  
 2401 tcaccacgga gaccatatcg accaccatgg agaacagtct caagtccggg aagggggcag  
 2461 ctgccatgat cccaggccca cagacggtgg ccacggaaat ccgttctctt tctccgatca  
 2521 tcgggaaaga tgtcctcacc agcacctacg gcgccactgc ggaaaccctc tcaacctcca  
 2581 ccaccacca tgtcaccaa actgtgaaag gaggggtttt tgagacaagg atcgagaagc  
 2641 gaatcatcat tactggggat gaagatgtcg atcaagacca ggccctgggt ttggccatca  
 2701 aggaggccaa actgcagcat cctgatatgc tggtaaccaa agctgtcgtat tacagagaaa  
 2761 cagacccatc cccagaggag agggacaaga agccacagga atcctgacct ctgtgaagag  
 2821 atcctggcat ttctggtcca acccaagcca gagaaccatt aagaaggggc cttcattctg  
 2881 gattctccga cgcaacactg acgtcccagc tgcgacgtac tgtcactgat gagagactgg  
 2941 gaagggaaaa gcatatatat atagatatat agagatatag atatataatc aggaaacacc  
 3001 gcatccttgc actgctgctg gggctggcag agcagttggc tgacagcaac aaccgacatc  
 3061 tgaacaccta catttccttt gcagacaaat tgaagaactg gtgggatttt tttcaagaaa  
 3121 aaaaattata taataactat aatcccttgc tcaccctttt cccccgcaa ataagaaacg  
 3181 caagccagac cacgatgatt gtagaagtcc ctcccgcctt ggttctgcac gttacagtta  
 3241 gcagacgagc aattccattt gttcttctcc agcatctcta aggccactt gaatgcaaag  
 3301 gaaaacactt gcacagcaaa gcaagagaag tcacagcagc aagacacgca cagtcaacca  
 3361 ttttccgaga aaaaaagaaa attccccact tggaaagaaa gaggaggaac actggattct  
 3421 tactttctgg atcttgacac tgggctgcaa aacctacctt cctctctccc gcctcccctc  
 3481 accctcaact ctcaatgtct tgctgtcatt ttctgtctcg gctccctcct ccccttccc  
 3541 ccttccccca cccacacccc ttcaccctct gtgtcctggt ccttctgagg gccactgcag  
 3601 atgactctcc tttgaaatga gaaaaagaaa agaaagcaag aacagaaaac gaagccacag  
 3661 gaagggaagt agacattgta tgcttatggt ttctcattat gaaggtgcag cttgtaggag  
 3721 gtttgtacgg atgtgctttg aagttatgta tattacatat aacaggaaaa aatattaata  
 3781 aacagtgctg gtaagtatga agctgacatt ctaaaattat aattatctga ctgtgattga  
 3841 tgtatcctga ggttcctaga tctcactgaa ctggcccagc taaggagacc tggactctgg  
 3901 gtgtggggtt gctcacagta ggggctgacg ggttcagtg agtaatactg tgtgtggtgt

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FIG. 32 (cont.)

3961 ttgtaattgg ttgattggtg gggaggggtg gggggcccta atggagaggt gtggggttgg  
 4021 caagaaagaa gcaacacaga tgtcgtcccc aaaatgccag ttcaagacac cttctccctg  
 4081 cccccctggt agtaacagtc agggcctggt ctgtgctcag gtactgggtc ccagtctggg  
 4141 actctgctgc tgaagtggcc acagtagagg tccctggctt agtccttata tccctacggg  
 4201 gcttgccttg gttttcagtc ttctctctct ttctctcttt tttttttttt tgccacattc  
 4261 tgcccttccc tgaccccat tgaataacca actccatata caaaggaggg tgggtgctctc  
 4321 agccattgta gaagatgggt gctttaacct gactgtctaa aaattcccag ctaagccttt  
 4381 tcctctactc tcttccttgt tctgaatcat ttcttcttct caggccaaag tagccatggt  
 4441 aaggaggctt catggggcag accctgaaag atcaaaaactg catttgcaaa gccctcccct  
 4501 gtcccaggac aaagctgaga ctgacgggtg atgttgctca taggctccag ctctgcataa  
 4561 gaccttggtt tggagacctc cctctcagtc aacagctgaa ctctgagctt gtgccagaa  
 4621 attaccccaa gaccacagga acccttcaag aagctcccat cacaagcttg gcattgctct  
 4681 ctgccacacg tgggcttcct caggcttgct tgccacaagc tacttctctg agctcagaaa  
 4741 gtgccccttg atgagggaaa atgtcccact gcactgcgaa tttctcagtt ccattttacc  
 4801 tcccagtcct ccttctaaac cagttaataa attcattcca caagtattta ctgattacct  
 4861 gcttggtgcca gggactattc tcaggctgaa gaagggtgga ggggaggggc gaacctgagg  
 4921 agccacctga gccagcttta tatttcaacc atggctggcc catctgagag catctcccca  
 4981 ctctcgccaa cctatcgagg catagcccag ggatgcccc aggcggccca ggttagatgc  
 5041 gtcccttttg cttgtcagtg atgacataca ccttagctgc ttagctgggt ctggcctgag  
 5101 gcagggcagg aaatcagaat agcatttgct tctctgggca aatgggaagt tcagcggggc  
 5161 agcagaatca gtggcattcc ccctggtgca ggccggtggg tccactccaa ctccccctga  
 5221 gtgtagcagc acactttcca tacaccaggt tctttctaca atcctgggtg aaaagccaca  
 5281 gaaccttctt cctgcccttc ttgagagttc cccctctttc tgggtcaaga gctggagtgg  
 5341 tggctccatc ctctctgggc cacttcggtc taggaactca tctttgcagg aaccaggagt  
 5401 cctgagcaca ctgaacacac ctcagaggga ggatccttgt tgtggatttt gcacctggct  
 5461 ttggggcagg ggtgaagtga ccaggcttag cttgtggagt ttatgggcca ccagggtttg  
 5521 gggaaatcac catcccggg atgctgtgac ctcccttcta cggagatgca ggcagtgcc  
 5581 cgagggagga ggggacctgc aaagctagaa tctagggcac tgtttcctcc ccactcttct  
 5641 cttttagtag aatagagacg tttgtcttgt ctgtcttcaa cctacttttc cttttctctt  
 5701 ttttgtttct catcctctct gtgccacctc tccaccagg aggccatgta gcatagtgga  
 5761 aaaagtccct gagggcggtt aggagttctg ggtgaccatc ctggctcagc tcctaactca  
 5821 ccatgtgaca tcaggctatc cccattcccc ctcttgggac tcagtttccc gacttgcaaa  
 5881 ataagcagaa agaaccagat gctctccagg gtctttttct actttgctat ctcatgggtc  
 5941 ttcattttct cttattttgt tttctctgga tcttttccat ctgagggtac aggaagtacc  
 6001 aggacctgtt tcagtttttg aatcctgcaa gcacattcca agactggcct gaaactgcat  
 6061 gagcaacatc actcgaaata attttttttt tcaaaagcac cttacaacc aattgcgatg  
 6121 ctgtcctggt cctttttact cacacccttc tctcttct cgtccccatg ctccccacc

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## FIG. 32 (cont.)

6181 tcagtgtcc gtgctgtatg cgtgtgctct ctgttcttgt atactcaata taagtgaat  
6241 aaatgtgttt gatgctgaac cat

## Translation:

SAGGGVAEQAAPQSPPRPRAAPPRLPARGAEGAAPRPTCPTWGTPGPGVLVTMTTET  
GPDSEVKKAQEEAPQQPEAAA VTPVT PAGHGHPEANSNEKHPSQQDTRPAEQSLDM  
EEKDYSEADGLSERTTPSKAQKSPQKI AKKYKSAICRVTL LDASEYECEVEKHGRGQV  
LFDLVCEHLN LLEKDYFGLT FCDADSQKNWLDPSKEIKKQIRSSPWNFAFTVKFYPPD  
PAQLTEDITRYYLCLQLRADIITGRLP CSFVTHALLGSYAVQAE LGDYDAEEHVGNV  
SELRFPNQ TRELEERIMELHKTYRGMT PGEAEIH FLEN AKKLSMYGVDLHHAKDSEG  
IDIMLGVCANGLLIYRDLRLINRFAWPKILKISYKRSNFYIKIRPGEYEQFESTIGFK  
LPNHRSAKRLWKVCI EHHTFFRLVSPEPPP KGLVMGSKFRYSGRTQAQTRQASALID  
RPAPFFERSSSKRYTMSRSLDGA EFSRPASVSENHDAGPDGDKRDE DGE SGGQRSEAE  
EGEVRTPTKIKELKPEQETTPRHKQEF LDKPEDVLLKHQASINELKRTLKEPNSKLIH  
RDRDWERERRLPSSPASPS PKGTPEKANERAGLREGSEEKVKPPRPAPESDTGDEDQ  
DQERDTVFLKDNHLAI ERKCSSITV SSTSSLEAEVDFTVIGDYHGSAFEDFSRSLPEL  
DRDKSDSDTEGLLFSRDLNKGAPSQDDESGGIEDSPDRGACSTPDMPQFEPVKTETMT  
VSSLAIRKKIEPEAVLQTRVSAMDNTQQVDGSASVGREFIATTPSITTETISTTMENS  
LKSGKGAAAMIPGPQTVATEIRSLSPIIGKDVLTSTYGATAETLSTSTTHVTKTVKG  
GFSETRIEKRIIITGDEDVDQDQALALAIKEAKLQHPDMLVTKAVVYRETDPSPEERD  
KKPQES

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## FIG. 33

Human non-muscle alpha-actinin mRNA, complete cds -  
the second non-muscle alpha-actinin isoform designated ACTN4 (actinin-4)

ORIGIN

1 gcgcgccggc ggctcgggca gaggggaggc agctgaggcg ggagcggaca ggctgggtggg  
61 cgagcgagag gcgcggaatg gtggactacc acgcggcgaa ccagtcgtac cagtacggcc  
121 ccagcagcgc ggcaatggct tggcgggcggg ggagcatggg cgactacatg gcccaggagg  
181 acgactggga ccgggacctg ctgctggacc cggcctggga gaagcagcag cgcaagacct  
241 tcacggcatg gagcaactcc cacctgcgga aggcaggcac acagatcgag aacattgatg  
301 aggacttccg agacgggctc aagctcatgc tgctcctgga ggtcatatca ggggagcggg  
361 tacctaagcc ggagcggggg aagatgagag tgcacaaaat caacaatgtg aacaaagcgc  
421 tggactttat tgccagcaaa gggatcaagc tggacttcca tcgggcagaa gagattgtgg  
481 acggcaacgc aaagatgacc ctgggaatga tctggacat catccttagg ttcgccatcc  
541 aggacatctc cgtggaagag acctcggcca aggaagggt ccttctctgg tgccagagaa  
601 agacagcccc atataagaac gtcaatgtgc agaacttcca catcagctgg aaggatgggc  
661 ttgccttcaa tgccctgatc caccggcaca gaccagagct gattgagtat gacaagctga  
721 ggaaggacga ccctgtcacc aacctgaaca atgccttcga agtggctgag aaatacctcg  
781 acatcccaa gatgctggat gcagaggaca tcgtgaacac ggcccggccc gacgagaagg  
841 ccataatgac ctatgtgtcc agcttctacc atgccttttc aggagcgag aaggctgaaa  
901 ctgaaactgc cgccaaccgg atctgtgaagg tgctggctgt caaccaagag aactgcagca  
961 cctcgatgga ggactacgag aagctggcca gcgacctct ggagtggatc cggcgacca  
1021 tcccctggct ggaggaccgt gtgccccaaa agactatcca ggagatgcag cagaagctgg  
1081 aggacttccg cgactaccgg cgtgtgcaca agccgccccaa ggtgcaggag aagtgccagc  
1141 tggagatcaa cttcaacagc gtgcagacca agctgcgcct cagcaaccgg cccgccttca  
1201 tgccctccga gggcaagatg gtctcggaca tcaacaatgg ctggcagcac ttggagcagg  
1261 ctgagaaggg ctacgaggag tggctgctga atgagattcg caggctggag cggctcgacc  
1321 acctggcaga gaagttccgg cagaaagcct ccatccacga ggcctggact gacgggaagg  
1381 aagccatgct gaagcaccgg gactacgaga cggccacact atcggacatc aaagccctca  
1441 ttcgcaagca cgaggccttc gagagcgacc tggctgcgca ccaggaccgc gtggagcaga  
1501 tcgccgcctc cgcccaggag ctcaacgagc tggattacta cgactcccac aatgtcaaca  
1561 cccggtgcca gaagatctgt gaccagtggg acgcccctcg ctctctgaca catagtcgca  
1621 gggaagccct ggagaaaaca gagaagcagc tggaggccat catcgaccag ctgcacctgg  
1681 aatacgccaa gcccgcggcc cccttcaaca actggatgga gagcgccatg gaggacctcc  
1741 aggacatggt catcgtccat accatcgagg agattgagg cctgatctca gcccatgacc  
1801 agttcaagtc caccctgccg gacgcccata gggagcgaga ggccatcctg catccacaag  
1861 gaggccagag gatcgtgag agcaaccaca tcaagctgtc gggcagcaac ccctacacca  
1921 ccgtcaccctc gcaaatcatc aactccaagt gggagaaggt gcagcagctg gtgccaaaac  
1981 gggaccatgc ctcctggag gagcagagca agcagcagca gtccaacgag cacctgcgcc  
2041 gccagttcgc cagccaggcc aatgttgtgg ggccctggat ccagaccaag atggaggaga  
2101 tcgcgatctc cattgagatg aacgggaccc tggaggacca gctgagccac ctgaagcagt  
2161 atgaacgcag catcgtggac tacaagccca acctggacct gctggagcag cagcaccagc  
2221 tcatccagga ggccctcatc ttcgacaaca agcacacca ctataccatg gagcacatcc  
2281 gcgtgggctg ggagcagctg ctaccacca ttgcccgcac catcaacgag gtggagaacc  
2341 agatccttac ccgcgacgcc aagggcata gccaggagca gatgcaggag ttccgggcgt  
2401 ccttcaacca cttcgacaag gatcatggcg gggcgctggg gcgaggagtt caaggcctgc  
2461 ctcatcagcc tgggctacga cgtggagaac gaccggcagg tgaggccgag ttcaaccgca  
2521 tcatgagcct ggtcgacccc aaccatagcg gccttggttac cttccaagcc ttcacgact  
2581 tcatgtcgcg ggagaccacc gacaccgaca cggctgacca ggtaatcact tccttcaagg

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## FIG. 33 (cont.)

2641 tcctagcagg ggacaagaac ttcatacacag ctgaggagct gcggagagag ctgccccccg  
 2701 accaggccga gtactgcatc gcccgcatgg cgccatacca gggccctgac ggcgtgcgcg  
 2761 gtgccctcga ctacaagtcc ttctccacgg ccttgtagtg cgagagcgac ctgtgaggcc  
 2821 ccagagacct gacccaacac ccccgacgcc tccaggagcc tggcagcccc acagtcccat  
 2881 tcctccactc tgtatctatg caaagcactc tctctgcagt ctccgggggtg ggtgggtggg  
 2941 cagggagggg ctggggcagg ctctctcctc tctctctttg tgggttggcc aggaggttcc  
 3001 cccgaccagg ttggggagag ttggggccag cgcttctggt ctggtaaata tgtatgatgt  
 3061 gttgtgcttt tttaaccaag gaggggcccag tggattccca cagcacaacc ggtcccttcc  
 3121 atgccctggg atgcctcacc acaccagggt ctcttccttt gctctgaggt cccttcaagg  
 3181 cctccccaat ccaggccaaa gccccatgtg ccttggtccag ggaactgcct gggccatgcg  
 3241 agggggccagc agagggcgcc accacctgac ggctgggacc caccagccc ctctccctc  
 3301 tctgctccag actcacttgc cattgccagg agatggcccc aacaagcacc ccgcttttgc  
 3361 agcagaggag ctgagttggc agaccgggc cccctgaacc gcaccccatc ccaccagccc  
 3421 cggccttgct ttgtctggcc tcacgtgtct cagattttct aagaaccaa aaa

## Translation:

MVDYHAANQSYQYGPSSAAMAWRRGSMGDYMAQEDDWRDLLLLDPAWEKQQRKTFTAW  
 SNSHLRKAGTQIENIDEDFRDGLKLMLLLEVISGERLPKPERGKMRVHKINNVNKALD  
 FIASKGIKLDHFHRAEEIVDGNAMTLGMIWTIILRFQDISVEETSAKEGLLLWCQR  
 KTAPYKNVNVQNFHISWKDGLAFNALIHRHRELIEYDKLRKDDPVTNLNNAFEVAEK  
 YLDIPKMLDAEDIVNTARPDEKAIMTYVSSFYHAFSGAQKAETETAANRICKVLAVNQ  
 ENCSTSMEDYEKLASDLLEWIRRTIPWLEDRVPOKTIQEMQQKLEDFRDYRRVHKPPK  
 VQEKQLEINFNSVQTKLRLSNRPAFMPSEGKMSDINNGWQHLEQAEKGYEEWLLNE  
 IRRLERLDHLAEKFRQKASIEAWTDGKEAMLKHRDYETATLSDIKALIRKHEAFESD  
 LAHQDRVEQIAASAQELNELDYDSDHNVNTRCQKICDQWDALGSLTHSRREALEKTE  
 KQLEAIIIDQLHLEYAKPAAPFNNWMESAMEDLQDMFIVHTIEEIEGLISAHDQFKSTL  
 PDADREREAILHPQGGQRIAESNHIKLSGSNPYTTVTPQIINSKWEKVQQLVPRDHA  
 LLEEQSKQQQSNEHLRQFASQANVVGPIQTKMEEIAISIEMNGTLEDQLSHLKQYE  
 RSIVDYKPNLDLLEQQHQLIQEALIFDNKHTNYTMEHIRVGWEQLLTTIARTINEVEN  
 QILTRDAKGISQEQMQEFRASFNFHDKDHGGALGRGVQGLPHQPLRRGERPAGEAEF  
 NRIMSLVDPNHSGLVTFQAFIDFMSRETTDTADQVITSFKVLADGNFITAEEELRR  
 ELPPDQAEYCIARMAPYQGPDGVRGALDYKSFSTALYGESDL

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## FIG. 34

Homo sapiens actinin, alpha 4 (ACTN4) mRNA

ORIGIN

```

1  cgcgggccgcg tgcacctacc acgcgggcgaa ccagtcgtac cagtacggcc ccagcagcgc
61  gggcaatggc gctggcggcg ggggcagcat gggcgactac atggcccagg aggacgactg
121  ggaccgggac ctgctgctgg acccggcctg ggagaagcag cagcgcaaga ccttcacggc
181  atggtgcaac tcccacctgc ggaaggcagg cacacagatc gagaacattg atgaggactt
241  ccgagacggg ctcaagctca tgctgctcct ggaggtcata tcaggggagc ggttacctaa
301  gccggagcgg gggaagatga gagtgcacaa aatcaacaat gtgaacaaag cgctggactt
361  tattgccagc aaaggcgtca agctggtctc catcggggca gaagagattg tggacggcaa
421  cgcaaagatg accctgggaa tgatctggac catcatcctt aggttcgcca tccaggacat
481  ctccgtggaa gagacctcgg ccaaggaagg gctccttctc tgggtgccaga gaaagacagc
541  cccgtataag aacgtcaatg tgcagaactt ccacatcagc tgggaaggatg gtcttgccctt
601  caatgccctg atccaccggc acagaccaga gctgattgag tatgacaagc tgaggaagga
661  cgaccctgtc accaacctga acaatgcctt cgaagtggct gagaaatacc tcgacatccc
721  caagatgctg gatgcagagg acatcgtgaa cacggcccgg cccgacgaga aggccataat
781  gacctatgtg tccagcttct accatgcctt ttcaggagcg cagaaggctg aaactgccgc
841  caaccggatc tgtaagggtg tggctgtcaa ccaagagaac gagcacctga tggaggacta
901  cgagaagctg gccagcgacc tcctggagtg gatccggcgc accatcccct ggctggagga
961  ccgtgtgccc caaaagacta tccaggagat gcagcagaag ctggaggact tccgcgacta
1021  ccggcgtgtg cacaagccgc ccaaggtgca ggagaagtgc cagctggaga tcaacttcaa
1081  cacgctgcag accaagctgc gcctcagcaa ccggcccggc ttcatgcctt ccgagggcaa
1141  gatggtctcg gacatcaaca atggctggca gcacttggag caggctgaga agggctacga
1201  ggagtggctg ctgaatgaga tccgcaggct ggagcggctc gaccacctgg cagagaagtt
1261  ccggcagaag gcctccatcc acgaggcctg gactgacggg aaggaagcca tgctgaagca
1321  ccgggactac gagacggcca cactatcgga catcaaagcc ctcatcgca agcacgaggc
1381  cttcgagagc gacctggctg cgcaccagga ccgctggag cagatcgccg ccattgccc
1441  ggagctcaac gagctggatt actacgactc ccacaatgtc aacaccgggt gccagaagat
1501  ctgtgaccag tgggacgccc tcggctctct gacacatagt cgcagggaag ccctggagaa
1561  aacagagaag cagctggagg ccatcgacca gctgcacctg gaatacgcca agcgcgcggc
1621  ccccttcaac aactggatgg agagcgccat ggaggacctc caggacatgt tcatcgtcca
1681  taccatcgag gagattgagg gcctgatctc agcccatgac cagttcaagt ccaccctgcc
1741  ggacgccgat agggagcgcg aggccatcct ggccatccac aaggaggccc agaggatcgc
1801  tgagagcaac cacatcaagc tgtcgggcag caaccctac accaccgtca ccccgcaaat

```

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FIG. 34 (cont.)

1861 catcaactcc aagtgggaga aggtgcagca gctggtgcca aaacgggacc atgccctcct  
 1921 ggaggagcag agcaagcagc agtccaacga gcacctgcgc cgccagttcg ccagccaggc  
 1981 caatgttgtg gggccctgga tccagaccaa gatggaggag atcgggcgca tctccattga  
 2041 gatgaacggg accctggagg accagctgag ccacctgaag cagtatgaac gcagcatcgt  
 2101 ggactacaag cccaacctgg acctgctgga gcagcagcac cagctcatcc aggaggccct  
 2161 catcttcgac aacaagcaca ccaactatac catggagcac atccgcgtgg gctgggagca  
 2221 gctgctcacc accattgccc gcaccatcaa cgaggtggag aaccagatcc tcaccgcga  
 2281 cgccaagggc atcagccagg agcagatgca ggagttccgg gcgtccttca accacttcga  
 2341 caaggatcat ggcggggcgc tggggcccga ggagttcaag gcctgcctca tcagcctggg  
 2401 ctacgacgtg gagaacgacc ggcagggtga ggccgagttc aaccgcatca tgagcctggt  
 2461 cgaccccaac catagcggcc ttgtgacctt ccaagccttc atcgacttca tgtcgcggga  
 2521 gaccaccgac acggacacgg ctgaccaggt catcgcttcc ttcaaggtct tagcagggga  
 2581 caagaacttc atcacagctg aggagctgag gagagagctg ccccccgaac aggcgagta  
 2641 ctgcatcgcc cgcattggcg cataccaggg ccctgacgcc gtgcccgggt ccctcgacta  
 2701 caagtccttc tccacggcct tgtatggcga gagcgacctg tgaggcccca gagacctgac  
 2761 ccaacacccc cgacggcctc caggaggggc ctgggcagcc ccacagtccc attcctccac  
 2821 tctgtatcta tgcaaacgac tctctgcagt cctccggggt ggggtgggtgg gca

## Translation:

MGDYMAQEDDWRDRLLLDPAWEKQQRKFTTAWCNSHLRKAGTQIENIDEDFRDGLKMLL  
 LEVISGERLPKPERGKMRVHKINNVNKALDFIASKGVKLVSIGAEIIVDGNAMTLGMIW  
 TIILRFAIQDISVEETSAKEGLLLWCQRKTAPYKNVNVQNFHISWKDGLAFNALIHRHRP  
 ELIEYDKLRKDDPVTNLNNAFEVAEKYLDIPKMLDAEDIVNTARPDEKAIMTYVSSFYHA  
 FSGAQKAETAANRICKVLAVNQENEHLMEDYEKLASDLLEWIRRTIPWLEDVRVPQKTIQE  
 MQOKLEDFRDYRRVHKPPKVQEKQLEINFNTLQTKLRLSNRPAFMPSEGKMSVDINNGW  
 QHLEQAEGYEEWLLNEIRRLERLDHLAEKFRQKASIHEAWTDGKEAMLKHRDYETATLS  
 DIKALIRKHEAFESDLAAHQDRVEQIAAIAQELNELDYYDSHNVNTRCQKICDQWDALGS  
 LTHSRREALEKTEKQLEAIDQLHLEYAKRAAPFNNWMESAMEDLQDMFIVHTIEEIEGLI  
 SAHDQFKSTLPDADREREAILAIHKEAQRIAESNHIKLSGSNPYTTVTPQIINSKWEKVQ  
 QLVPKRDLALLEEQSKQSQSNEHLRRQFASQANVVGPIQTKMEEIGRISIEMNGTLEDQL  
 SHLKQYERSIVDYKPNLDLLEQQHQLIQEALIFDNKHTNYTMEHIRVGWEQLLTTIARTI  
 NEVENQILTRDAKGISQEQMQEFRAFNFHFDKDHGGALGPEEFKACLI SLGYDVENDROG  
 EAEFNRIMSLVDPNHSGLVTFQAFIDFMSRETTDTDTADQVIASFVLAGDKNFITAEEL  
 RRELPPDQAEYCIARMAPYQGPDAVPGALDYKSFSTALYGESDL

008760-0964960

## FIG. 35

## CLATHRIN COAT ASSEMBLY PROTEIN AP50

## ORIGIN

1 cagggtctggt ctcagagcga tgggcccgcag agactgatct gccgccatga ttggaggcct  
 61 attcatctat aatcacaagg gggagggtgct catctcccga gtctaccgag atgacatcgg  
 121 gaggaacgca gtggatgcct ttcgggtcaa tggtatccat gcccggcagc aggtgcgcag  
 181 ccccgctcacc aacattgctc gcaccagctt cttccacggt aagcgggtcca acatttggct  
 241 ggcagcagtc accaagcaga atgtcaacgc tgccatggtc ttcgaattcc tctataagat  
 301 gtgtgacgtg atggccgctt actttggcaa gatcagcgag gaaaacatca agaacaattt  
 361 tttgctcata tatgagctgc tggatgagat tctagacttt ggctaccac agaattccga  
 421 gacaggcgcg ctgaaaacct tcatcacgca gcagggcac aagagtcagc atcagacaaa  
 481 agaagagcag tcacagatca ccagccaggt aactgggcag attggctggc ggcgagaggg  
 541 catcaagtat cgtcggaatg agctcttcct ggatgtgctg gagagtgtga acctgtctcat  
 601 gtccccacaa gggcagggtgc tgagtgccca tgtgtcgggc cgggtgggtga tgaagagcta  
 661 cctgagtggc atgcctgaat gcaagtttgg gatgaatgac aagattgtta ttgaaaagca  
 721 gggcaaaggc acagctgatg aaacaagcaa gagcgggaag caatcaattg ccattgatga  
 781 ctgcaccttc caccagtgtg tgcgactcag caagtttgac tctgaacgca gcatcagctt  
 841 tatcccgcca gatggagagt ttgagcttat gaggtatcgc acaaccaagg acatcatcct  
 901 tcccttccgg gtgatccgc tagtgcgaga agtgggacgc accaaactgg aggtcaaggt  
 961 ggtcatcaag tccaacttta aacctcact gctggctcag aagattgagg tgaggatccc  
 1021 aacccccactg aacacaagcg ggggtgcaggt gatctgcatg aaggggaagg ccaagtacaa  
 1081 ggccagcgag aatgccatcg tgtggaagat caagcgcatg gcaggcatga aggaatcgca  
 1141 gatcagcgca gagattgagc ttctgcctac caacgacaag aagaaatggg ctcgaccccc  
 1201 cattedcatg aactttgagg tgccattcgc gccctctggc ctcaagggtgc gctacttgaa  
 1261 ggtgtttgaa ccgaagctga actacagcga ccatgatgtc atcaaatggg tgcgctacat  
 1321 tggccgcagt ggcatttatg aaactcgtg ctagtgcga ctaggcagct agcccacctc  
 1381 cccagccacc ctctccaca ggtccaggtg ccgctccctc ccccaccaca catcagtgtc  
 1441 tcctccctcc tgctttgctg ccttcccttt gcaccagccc gagtctaggt ctggggccaag  
 1501 cacattacaa gtgggaccgg tggagcagcc cctgggctcc ctgggagggg gagttctgag  
 1561 gctcctgctc tcccatccac ctgtctgtcc tggcctaatt ccaggctctg agttctgtga  
 1621 ccaaagccag gtgggttccc tttccttccc acccctgtgg ccacagctct ggagtgggag  
 1681 gggttggtgc cctcacctc agagctcccc caaaggccag taatggatcc ccggcctcag  
 1741 tccctactct gctttgggat agtgtgagct tcattttgta cacgtgttgc ttcgtccagt  
 1801 taaaaaccca ataaactctg tagagtgg

## Translation:

MIGGLFIYNHKGEVLI SRVYRDDIGRNAVDAFRVNVIHARQQVRSPVTNIARTSFFHV  
 KRSNIWLA AVTKQNVNAAMVFEFLYKMC DVMAAYFGKISEENIKNNFLLIYELLDEIL  
 DFGYPQNS ETGALKTFITQQGIKSQHQTKEEQS QITSQVTGQIGWRREGIKYRRNELF  
 LDVLESVNLLMS PQGQVLSAHVSGRVVMKSYLSGMPECKFGMNDKIVIEKQKGKTADE  
 TSKSGKQSIAID DCTFHQCVRLSKFD SERSISFI PP DGEFELMRYRTTKDIILPFRVI  
 PLVREVGRTKLEVKVVIKSNFKPSLLAQKIEVRIPTPLNTSGVQVICMKGKAKYKASE  
 NAI VWKIKRMAGMKESQISAEIELLP TNDKKKWARPPISMNFEVPFAPSGLKVRYLKV  
 FEPKLNYS DHDVIKVVRYIGRS GIYETRC

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## FIG. 36

Homo sapiens GLUT1 C-terminal binding protein (GLUT1CBP) mRNA

ORIGIN

1 cacgggggagg cggaggcagc ggcggcgggcg gcggcgggcg cgggcgggcg ggagcagatc  
 61 ttctggtgac ccacttctc gctgctcatg ccgctgggac tggggcgccg gaaaaaggcg  
 121 cccctctag tggaaaatga ggaggctgag ccaggccgtg gagggctggg cgtgggggag  
 181 ccagggcctt tgggcggagg tgggtcgggg ggcccccaa tgggcttgcc cccctccc  
 241 ccagccctgc ggccccgcct tgtgttccac acccagctgg cccatggcag tccactggc  
 301 cgcacgagg ggttcaccaa cgtcaaggag ctgtatggca agattgccga ggccttccgc  
 361 ctgccaactg ccgaggtgat gttttgcacc ctgaacaccc acaaagtga catggacaag  
 421 ctctggggg gccaatcgg gctggaggac ttcattctcg cccacgtgaa ggggcagcgc  
 481 aaggaggtg aggtgttcaa gtcggaggat gcaactcggc tcaccatcac ggacaacggg  
 541 gctggctacg ccttcatcaa gcgcatcaag gagggcagcg tgatcgacca catccacctc  
 601 atcagcgtg gcgacatgat cgaggccatt aacgggcaga gcctgctggg ctgccggcac  
 661 tacgaagtg cccggctgct caaggaactg ccccgaggcc gtaccttcac gctgaagtc  
 721 acggagcctc gcaaggcctt cgacatgatc agccagcgtt cagcgggtg ccgccctggc  
 781 tctggccac aactgggcac tggccgagg accctgcggc tccgatcccg gggccccgcc  
 841 acggtggagg atctgccctc tgcctttgaa gagaaggcca ttgagaaggt ggatgacctg  
 901 ctggagagtt acatgggtat cagggacacg gagctggcgg ccaccatggt ggagctggga  
 961 aaggacaaaa ggaaccggga tgagctggcc gaggccctgg acgaacggct gggtgacttt  
 1021 gccttccctg acgagttcgt ctttgacgtc tggggcgcca ttggggacgc caaggtcggc  
 1081 cgctactagg actgcccccg gaccctgcga tgatgaccgg ggcgcaacct ggtggggggc  
 1141 cccagcaggg aactgacgt caggaccgga gcctccaagc ctgagcctag ctgagcagcc  
 1201 caaggacgat ggtgagggga ggtggggcca ggccccctgc cccgctccaa tcggtaccat  
 1261 cccctccctg gttcccagtc tggccggggg ccccgcccc cctgtgccct gttccccacc  
 1321 ctacctcagc tggggtcagg cacagggaag gggagggatc agccaaattt gggcgccac  
 1381 cccgcctcc accactttcc accatcagct gccaaactgg tccctctgtc tccctggggc  
 1441 cttgggttct gtttgggggt catgaccttc ctagtctcct gacgcaggga atacagggga  
 1501 gaggggtgtc cttcccccca gcaaatgcaa taatgccctc acccctcctg agaggagccc  
 1561 cctccctgtg gagcctgtta cctccgcatt tgacacgagt tgctgtgaac cccgcaacct  
 1621 cctccccacc tcccatctct ccttcaggc ccacccctgg cccagagcag gagggaggga  
 1681 gggacgatgg cggtgggttt ttgtatctga atttgctgtc ttgaacataa agaattctatc  
 1741 tgctgttaaa aaaaaaaaaa aaaaa

Translation:

MPLGLGRRKKAPPLVENEAEPRGGLGVGEPGPLGGGSGGPQMGLPPPPPALRPRL  
 VFHTQLAHGSPTGRIEGFTNVKELYGKIAEAFRLPTAEVMFCTLNTHKVDMDKLLGGQ  
 IGLEDFI FAHVKGQRKEVEVFKSEDALGLTITDNGAGYAFIKRIKEGSVIDHIHLISV  
 GDMIEAINGQSLLGCRHYEVARLLKELPRGRTFTLKLTEPRKAFDMISQRSAGGRPGS  
 GPQLGTGRGTLRLRSRGPATVEDLPFAFEKAIEKVDDLLESYMGIRDTELAATMVVEL  
 GKDKRNPDELAELDERLGDFAFPDEFVFDVWGAIGDAKVGRY

## FIG. 37

gp130 associated protein GAM

## ORIGIN

1 ggccgcccgg cgcccccagc agnccgagcc ggggcgcaca gncggggngc agaccgcgcc  
 61 ccccgccgcg attgacatga tgtttccaca aagcaggcat tcgggctcct cgcacctacc  
 121 ccagcaactc aaattcacca cctcggactc ctgcgaccgc atcaaagacg aatttcagct  
 181 actgc'aagct cagtaccaca gcctcaagct cgaatgtgac aagttggcca gtgagaagtc  
 241 agagatgcag cgtcactatg tgatgtacta cgagatgtcc tacggcttga acatcgagat  
 301 gcacaaacag gctgagatcg tcaaaaggct gaacgggatt tgtgccagg tcctgcccta  
 361 cctctcccaa gagcaccagc agcaggctctt gggagccatt gagagggcca agcagggtcac  
 421 cgctcccagag ctgaactcta tcatccgaca gcagctccaa gcccaccagc tgtcccagct  
 481 gcaggccctg gccctgccct tgacccact acccgtgggg ctgcagccgc cttcgctgcc  
 541 ggcggtcagc gcaggcaccg gcctcctctc gctgtccgcg ctgggttccc aggccacct  
 601 ctccaaggaa gacaagaacg ggcacgatgg tgacaccac caggaggatg atggcgagaa  
 661 gtcggattag cagggggccg ggacggggag gttgggagg gggacagagg ggagacagag  
 721 gcacggagag aaaggaatgt ttagcacaag acacagcgga gctcgggatg ggctaaactc  
 781 ccatagtatt tatggtggcc gccggcgggg gccccagccc agcttgagg ccacctctag  
 841 ctttcttccc taccocatcc ccggcttccc tcctcctccc tgcagcctgg ttaggtggat  
 901 acctgccctg acatgtgagg caagctaagg cctggaggga cagctgggag accaggtccc  
 961 aaggagcaa gacctcgca agcgagcag acccgccct ttccccgtt taggcatgtg  
 1021 taaccgacag tctgcctggg ccacagccct ctcaacctgg tactgcatgc acgcaatgct  
 1081 agctgcccct ttcccgtcct ggnaccccg agtctcccc gaccccggt cccaggtatg  
 1141 ctcccacctc cacctgcccc actcaccacc tctgctagtt ccagacacct ccacgcccac  
 1201 ctggtcctct cctaccgcac acaaaagggg gggaacgagg gacgagctta gctgagctgg  
 1261 gaggagcagg gtgagggtgg gcgaccagg attccccctc cccttcccaa ataacc

## Translation:

MFPQSRHSGSSSLPQQLKFTTSDSCDRIKDEFQLLQAQYHSLKLECDKLASEKSEMQR  
 HYVMYYEMSYGLNIEMHKQAEIVKRLNGICAQVLPYLSQEHQQQVLGAIERAKQVTAP  
 ELNSII RQQLQAHQLSQLQALALPLTPLPVGLQPPSLPAVSAGTGLLSLSALGSQAHL  
 SKEDKNHGDGTHQEDDGEKSD

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## FIG. 38

Homo sapiens amino-terminal enhancer of split (AES) mRNA

ORIGIN

```

1  ggccgcccgg cgccccagc agnccgagcc ggggcgcaca gncggggcgc agccccgcgc
61  ccccgccgcg attgacatga tgtttccaca aagcaggcat tcgggctcct cgcacctacc
121 ccagcaactc aaattcacca cctcggaactc ctgcgaccgc atcaaagacg aatttcagct
181 actgcaagct cagtaccaca gcctcaagct cgaatgtgac aagttggcca gtgagaagtc
241 agagatgcag cgtcactatg tgatgtacta cgagatgtcc tacggcttga acatcgagat
301 gcacaaacag gctgagatcg tcaaaaggct gaacgggatt tgtgcccagg tcctgccccta
361 cctctcccaa gagcaccagc agcaggctctt gggagccatt gagagggcca agcagggtcac
421 cgctcccagag ctgaactcta tcatccgaca gcagctccaa gccaccagc tgtcccagct
481 gcaggccctg gccctgccct tgaccccact acccgtaggg ctgcagccgc cttcgtgccc
541 ggcggtcagc gcaggcaccg gcctcctctc gctgtccgcg ctgggttccc agggccacct
601 ctccaaggaa gacaagaacg ggcacgatgg tgacaccac caggaggatg atggcgagaa
661 gtcggattag cagggggccg ggacaggag gttgggaggg gggacagagg ggagacagag
721 gcacggagag aaaggaatgt ttagcacaag acacagcgga gctcgggatt ggctaattctc
781 ccatagtatt tatggtggcg ccggcggggc cccagcccag cttgcaggcc acctctagct
841 ttcttcttac cccattccgg cttccctcct cctcccctgc agcctggta ggtggatacc
901 tgccctgaca tgtgaggcaa gctaaggcct ggagggtcag atgggagacc aggtcccaag
961 ggagcaagac ctgcgaagcg cagcagcccc ggcccttccc ccgttttgaa catgtgtaac
1021 cgacagtctg ccctgggcca cagccctctc accctggta tgcatgcacg caatgctagc
1081 tgcccttttc ccgtcctggg caccctcagt ctcccctgac cccgggtccc aggtatgctc
1141 ccacctccac ctgccccact caccacctct gctagtcca gacacctcca cgcccacctg
1201 gtcctctccc atcgcccaca aaaggggggg cacgaggac gagcttagct gagctgggag
1261 gagcagggtg agggtagggc acccaggatt cccctcccc ttcccaata aagatgaggg
1321 tact

```

Translation:

```

MMFPQSRHSGSSHLPPQLKFTTSDSCDRIKDEFQLLOAQYHSLKLECDKLASEKSEMQ
RHYVVMYYEMSYGLNIEMHKQAEIVKRLNGICAQVLPYLSQEHQQQVLGAIERAKQVTA
PELNSIIRQQLQAHQLSQLQALALPLTLPVGLQPPSLPAVSAGTGLLSLSALGSQAH
LSKEDKNGHDGDTHQEDDGEKSD

```

FIG. 39

Antiquitin 1 (antiquitin=26g turgor protein homolog), mRNA

## ORIGIN

1 cctgctccaa ggtccagaga gctttctggt ctttgcagca ggcctgccgc cttcatgtcc  
 61 actctcctca tcaatcagcc ccagtatgcg tggctgaaag agctggggct ccgcgaggaa  
 121 aacgagggcg tgtataatgg aagctgggga ggccggggag aggttattac gacctattgc  
 181 cccgctaaca acgagccaat agcaagagtc cgacaggcca gtgtggcaga ctatgaagaa  
 241 actgtaaaga aagcaagaga agcatggaaa atctgggcag atattcctgc tccaaaacga  
 301 ggagaaatag taagacagat tggcgatgcc ttgctgggaga agatccaagt actaggaagc  
 361 ttggtgtctt tggagatggg gaaaatctta gtggaagggt tgggtgaagt tcaggagtat  
 421 gtggatatct gtgactatgc tgttggttta tcaaggatga ttggaggacc tatcttgccct  
 481 tctgaaagat ctggccatgc actgattgag cagtggaaac ccgtaggcct ggttggaaac  
 541 atcacggcat tcaatttccc tgtggcagtg tatggttga acaacgccat cgccatgac  
 601 tgtggaaatg tctgcctctg gaaaggagct ccaaccactt ccctcattag tgtggctgtc  
 661 acaaagataa tagccaaggt tctggaggac aacaagctgc ctggtgcaat ttgttccttg  
 721 acttggtgtg gagcagatat tggcacagca atggccaaag atgaacgagt gaacctgctg  
 781 tccttcactg ggagcactca ggtgggaaaa cagggtgggc tgatggtgca ggagaggttt  
 841 gggagaagtc tgttggaact tggaggaaac aatgccatta ttgcctttga agatgcagac  
 901 ctcagcttag ttgttccatc agctctcttc gctgctgtgg gaacagctgg ccagaggtgt  
 961 accactgcca ggcgactgtt tatacatgaa agcatccatg atgaggttgt aaacagactt  
 1021 aaaaaggcct atgcacagat ccgagttggg aacctatggg accctaattg tctctatggg  
 1081 ccactccaca ccaagcaggc agtgagcatg tttcttggag cagtggaaaga agcaaagaaa  
 1141 gaaggtggca cagtgttcta tgggggcaag gttatggatc gccctggaaa ttatgtagaa  
 1201 ccgacaattg tgacaggtct tggccacgat gcgtccattg cacacacaga gactttcgct  
 1261 ccgattctct atgtctttaa attcaagaat gaagaagagg tctttgcatg gaataatgaa  
 1321 gtaaaacagg gactttcaag tagcatcttt accaaagatc tgggcagaat ctttcgctgg  
 1381 cttggacctt aaggatcaga ctgtggcatt gtaaatgtca acattccaac aagtggggct  
 1441 gagattggag gtgccttttg aggagaaaag cacactggtg gtggcagga gtctggcagt  
 1501 gatgcctgga aacagtacat gagaaggctt acttgacta tcaactacag taaagacctt  
 1561 cctctggccc aaggaatcaa gtttcagtaa aggtgtttta gatgaacatc ccttaattg  
 1621 aggtgttcca gcagctgttt ttggagaaga caaagaagat taaagttttc cctgaataaa  
 1681 tgcattatta tgactgtgac agtgactaat cccctatga ccccaaagcc ctgattaaat  
 1741 caagagattc cttttttaa aatcaaaata aaattgttac aacatagcca tagttactaa  
 1801 aaaaaaaaa

## Translation:

MSTLLINQPQYAWLKELGLREENEGVYNGSWGGRGEVITTYCPANNEPIARVRQASVA  
 DYEETVKKAREAWKIWADI PAPKRGEIVRQIGDALREKIQVLGSLVSLMGKILVEGV  
 GEVQEYVDICDYAVGLSRMIGGPILPSERSGHALIEQWNPVGLVGIITAFNFPVAVYG  
 WNNAIAMICGNVCLWKGAPTTSLISVAVTKIIAKVLEDNKLPGAICSLTCGGADIGTA  
 MAKDERVNLLSFTGSTQVGKQVGLMVQERFGRSLLELGGNNAI IAFEDADLSLVVPSA  
 LFAAVGTAGQRCTTARRLFIHESI HDEVVNRLKKAYAQIRVGNPWDPNVLYGPLHTKQ  
 AVSMFLGAVEEAKKEGGTVVYGGKVMDRPGNYVEPTIVTGLGHDASIAHTETFPILY  
 VFKFKNEEEVFAWNNEVKQGLSSSI FTKDLGRI FRWLGPKGSDCGIVNVNIPTSGAEI  
 GGAFGGEKHTGGGRESGSDAWKQYMRRSTCTINYSKDLPLAQGIKFQ



49/52

# FIG. 40

ARP2/3 protein COMPLEX 41 KD SUBUNIT (P41-ARC), mRNA

ORIGIN

```

1 ggcacgaggg agcccagagc cggttcggcg cgtcgactgc ccagagtccg cggccggggc
61 gcgggaggag ccaagccgcc atggcctacc acagcttcct ggtggagccc atcagctgcc
121 acgcctggaa caaggaccgc acccagattg ccatctgccc caacaaccat gaggtgcata
181 tctatgaaaa gagcgggtgcc aaatggacca aggtgcacga gctcaaggag cacaacgggc
241 aggtgacagg catcgactgg gcccccgaga gtaaccgtat tgtgacctgc ggcacagacc
301 gcaacgccta cgtgtggacg ctgaagggcc gcacatggaa gcccacgctg gtcatcctgc
361 ggatcaaccg ggctgcccgc tgcgtgcgct gggcccccaa cgagaacaag tttgctgtgg
421 gcagcggctc tcgtgtgata tccatctgtt atttcgagca ggagaatgac tgggtgggtt
481 gcaagcacat caagaagccc atccgctcca ccgtcctcag cctggactgg caccccaaca
541 atgtgctgct ggctgccggc tcctgtgact tcaagtgtcg gatcttttca gcctacatca
601 aggaggtgga ggaacggccg gcacccaccc cgtggggctc caagatgccc tttggggaac
661 tgatgttcga atccagcagt agctgcggtt gggatcatgg cgtctgtttc tcagccagcg
721 ggagccgcgt ggccctgggtg agccacgaca gcaccgtctg cctggctgat gccgacaaga
781 agatggccgt cgcgactctg gcctctgaaa cactaccact gctggcgctg accttcatca
841 cagacaacag cctggtggca gcggggccacg actgcttccc ggtgctgttc acctatgacg
901 ccgccgcggg gatgctgagc ttcggcgggc ggctggacgt tcctaagcag agctcgcagc
961 gtggccttgac ggcccgcgag cgcttccaga acctggacaa gaaggcgagc tccgaggggtg
1021 gcacggctgc gggcgcgggc ctagactcgc tgcacaagaa cagcgtcagc cagatctcgg
1081 tgctcagcgg cggcaaggcc aagtgtcgc agttctgcac cactggcatg gatggcggca
1141 tgagtatctg ggatgtgaag agcttggagt cagccttgaa ggacctcaag atcaaataac
1201 ctgtgaggaa tatgttgctt tcacctaacc tgctggggaa gcggggagag gggtcaggga
1261 ggctaattgt tgctttgctg aatgtttctg gggtagcaat acgagttccc ataggggctg
1321 ctccctcaaa aaggaggagg acagatgggg agcttttctt acctattcaa ggaatacgtg
1381 cctttttctt aaatgctttc atttattgaa aaaaaaaaaa aaaaaaaa

```

Translation:

MAYHSFLVEPISCHAWNKDRTQIAICPNNHEVHIYEKSGAKWTKVHELKEHNGQVTGI  
DWAPESNRIVTCGTDRNAYVWTLKGRWTKPTLVILRINRAARCVRWAPNENKFAVGSG  
SRVISICYFEQENDWWVCKHIKKPIRSTVLSLDWHPNNVLLAAGSCDFKCRI FSAYIK  
EVEERPAPTPWGSKMPFGELMFESSSSCGWVHGVCFSASGSRVAWVSHDSTVCLADAD  
KKMAVATLASETLPLLALTFITDNSLVAAGHDCFVLFYDAAAGMLSFGGRLDVPKQ  
SSQRLTARERFQNLDDKASSEGGTAAGAGLDSLHKNSVSQISVLSGGKAKCSQFCTT  
GMDGGMSIWDVKSLESALKDLKIK

008160-8564950

## FIG. 41a

H.sapiens seb4D mRNA

ORIGIN

1 gagcgcgggt ttctcgcggc ccctggccgc ccccggcgtc atgtacggct cgcagaaggg  
 61 caccacgttc accaagatct tcgtgggagg cctgccgtac cacactaccg acgcctcgct  
 121 caggaagtac ttcgagggct tcggcgacat cgaggaggcc gtgggtcatca ccgaccgcca  
 181 gacgggcaag tcccgcggct acggcttcgt gaccatggcc gaccgggcgg cagctgagag  
 241 ggcttgcaaa gaccctaacc ccatcatcga cggccgcaag gccaacgtga acctggcata  
 301 tctgggcgcc aagccttggt gtctccagac gggctttgcc attggcgtgc agcagctgca  
 361 ccccaccttg atccagcgga cttacgggct gaccccgcac tacatctacc caccagccat  
 421 cgtgcagccc agcgtggtga tccagccgc ccctgtcccg tcgctgtcct cgccctacat  
 481 tgagtacacg ccggccagcc cggctctacg ccagtaccca ccggccacct atgaccagta  
 541 cccatacgcc gcctcgcctg ccacggctga cagcttcgtg ggctacagct accctgccgc  
 601 cgtgcaccag gccctctcag ccgcagcacc cgcgggcacc actttcgtgc agtaccaggg  
 661 gccgcagctg cagcctgaca ggatgcagtg aggggcgttc ctgccccgag gactgtggca  
 721 ttgtcacctt cacagcagac agagctgcca ggccatgatg ggctggcgac agccgggctg  
 781 agcttcagtg aggtgccacc agcaccctg cctccgaaga ccgctcgggc attccgcctg  
 841 cgccctggga cagcggagag acggcttctc tttaatctag gtcccattgt gtcttgaggg  
 901 aggactttta agaatgactg agaactattt aaagacgcaa tcccaggttc cttgcacacc  
 961 atggcagcct ctctctgcac cttctcctgc ctctccacac tccaggttcc ctcaggcttg  
 1021 tgtccccact gctgcatcgt ggccggggtg cacagaccct ctgcagcccc tggctgccct  
 1081 ggactgtgca gagatgcctg actccaggga aacctgaaag caagaagtta atggactgtt  
 1141 tattgtaact tgatcctccc gagctgtgag cgcagtctga ggtctgagga caggcctcc  
 1201 tgttgagtc ccattttctc catcagggca cgtgggcggc ttcctcaagc ccggaggagc  
 1261 tcccaggcgc acaggggccc ccggtaacag gggccgcccg ccaaaggccc ctttccagtc  
 1321 atagcactga agttgcaact tttttcttgt aattgtttg ctactaagat aatttcagaa  
 1381 gttcagtcta ttttttcagc ggatactgcc gccaccaaga atccaaacct aggaa

Translation:

SAGFSRPLAAPGVMYGSQKGTTFTKIFVGGLPYHTTDASLRKYFEGFGDIEEAVVITD  
 RQTGKSRGYGFVTMADRAAAERACKDPNPIIDGRKANVNLAYLGAKPWCLQTGFAIGV  
 QQLHPTLIQRTYGLTPHYIYPPAIVQPSVVI PAAPVPSLSSPYIEYTPASPVYAQYPP  
 ATYDQYPYAASPATADSFVGYSYPAAVHQALSAAAPAGTTFVQYQAPQLQPDRLQ

0064958-091800

FIG. 41b

H.sapiens seb4B mRNA

ORIGIN

1 gcggcggatg cagtacaacc ggcgctttgt caacgttgtg cccacctttg gcaagaagaa  
 61 gggcaccacg ttcaccaaga tcttcgtggg cggcctgccg taccacacta ccgacgcctc  
 121 gctcaggaag tacttcgagg gcttcggcga catcgaggag gccgtgggtca tcaccgaccg  
 181 ccagacgggc aagtcccgcg gctacggctt cgtgaccatg gccgaccggg cggcagctga  
 241 gagggcttgc aaagacccta accccatcat cgacggccgc aaggccaacg tgaacctggc  
 301 atatctgggc gccaaacctt ggtgtctcca gacgggcttt gccattggcg tgcagcagct  
 361 gacccccacc ttgatccagc ggacttacgg gctgaccccg cactacatct acccaccagc  
 421 catcgtgcag cccagcgtgg tgatcccagc cgcccctgtc ccgtcgtgtgt cctcgcctta  
 481 cattgagtag acgccggcca gcccggtcta cgcccagtag ccaccggcca cctatgacca  
 541 gtaccatac gccgcctcgc ctgccacggc tgacagcttc gtgggctaca gctaccctgc  
 601 cgccgtgcac caggccctct cagccgcagc acccgcgggc accactttcg tgcagtacca  
 661 ggcgccgcag ctgcagcctg acaggatgca gtgaggggcg ttcctgcccc gaggactgtg  
 721 gcattgtcac cttcacagca gacagagctg ccaggccatg atgggctggc gacagccccg  
 781 ctgagcttca gtgaggtgcc accagcaccg gtgcctccga agaccgctcg ggcattccgc  
 841 ctgcgccctg ggacagcgga gagacggctt ctctttaatc taggtcccat tgtgtcttga  
 901 gggaggactt ttaagaatga ctgagaacta tttaaagacg caatcccagg ttccttgcac  
 961 accatggcag cctctccttg cactttctcc tgcctctcca cactccaggt tccctcaggc  
 1021 ttgtgtcccc actgctgcat cgtggcgggg tgtcacagac cctctgcagc ccctggctgc  
 1081 cctggactgt gcagagatgc ctgactccag ggaaacctga aagcaagaag ttaatggact  
 1141 gtttattgta acttgatcct cccgagctgt gagcgagtc tgaggctgta ggacacggcc  
 1201 tcctgttgga gtcccatttt ctccatcagg gcacgtgggc ggcttcctca agcccggagg  
 1261 agctcccagg cgacacgggg ccgccggtaa caggggccgc cggccaaagg cccctttcca  
 1321 gtcatagcac tgaagttgca acttttttct tgtaattgtt ttgctactaa gataatttca  
 1381 gaagttcagt ctattttttc agcggatact gccgccacca agaatccaaa cctaggaa

Translation:

RRMQYNRRFVNVPVTFGKKKGTTFTKIFVGGLPYHTTDASLRKYFEGFGDIEEAVVIT  
 DRQTGKSRGYGFVTMADRAAAERACKDPNPIIDGRKANVNLAYLGAKPWCLQTGFAIG  
 VQQLHPTLIQRTYGLTPHYIYPPAIVQPSVVI PAAPVPSLSPPYIEYTPASPVYAQYP  
 PATYDQYPYAASPATADSFVGYSYPAAVHQALSAAAPAGTTFVQYQAPQLQPD RMQ

FIG. 42

Homo sapiens lamin A/C (LMNA) mRNA

ORIGIN

1 actcagtgtt cgcgggagcc gcacctacac cagccaaccc agatccccgag gtccgacagc  
 61 gcccggccca gatccccacg cctgccagga gcaagccgag agccagccgg ccggcgccact  
 121 ccgactccga gcagtctctg tccttcgacc cgagccccgc gccctttccg ggacccttgc  
 181 cccgcgggca gcgctgccaa cctgccggcc atggagaccc cgtcccagcg gcgcgccacc  
 241 cgagcgggg cgagggccag ctccactccg ctgtcgccca cccgcatcac ccggctgcag  
 301 gagaaggagg acctgcagga gctcaatgat cgcttgccgg tctacatcga ccgtgtgcgc  
 361 tcgctggaaa cggagaacgc agggctgcgc cttcgcatca ccgagtctga agaggtggtc  
 421 agccgcgagg tgtccggcat caaggccgcc tacgaggccg agctcgggga tgcccgcgaag  
 481 acccttgact cagtagccaa ggagcgcgcc cgcctgcagc tggagctgag caaagtgcgt  
 541 gaggagttaa aggagctgaa agcgcgcaat accaagaagg agggtgacct gatagctgct  
 601 caggctcggc tgaaggacct ggaggctctg ctgaactcca aggaggccgc actgagcact  
 661 gctctcagtg agaagcgac gctggagggc gagctgcatg atctgcgggg ccaggtggcc  
 721 aagcttgagg cagccctagg tgaggccaag aagcaacttc aggatgagat gctgcggcgg  
 781 gtggatgctg agaacaggct gcagaccatg aaggaggaac tggacttcca gaagaacatc  
 841 tacagtgagg agctgcgtga gaccaagcgc cgtcatgaga cccgactggt ggagattgac  
 901 aatgggaagc agcgtgagtt tgagagccgg ctggcggatg cgctgcagga actgcgggcc  
 961 cagcatgagg accaggtgga gcagtataag aaggagctgg agaagactta ttctgccaa  
 1021 ctggacaatg ccaggcagtc tgctgagagg aacagcaacc tgggtgggggc tgcccacgag  
 1081 gagctgcagc agtcgcgcat ccgcatcgac agcctctctg cccagctcag ccagctccag  
 1141 aagcagctgg cagccaagga ggcgaagctt cgagacctgg aggactcact ggcccgtagg  
 1201 cgggacacca gccggcggct gctggcggaa aaggagcggg agatggccga gatgcgggca  
 1261 aggatgcagc agcagctgga cgagtaccag gagcttctgg acatcaagct ggccctggac  
 1321 atggagatcc acgcctaccg caagctcttg gagggcgagg aggagaggct acgcctgtcc  
 1381 cccagcccta cctcgcagcg cagccgtggc cgtgcttcct ctactcatc ccagacacag  
 1441 ggtgggggca gcgtcaccaa aaagcgcaaa ctggagtcca ctgagagccg cagcagcttc  
 1501 tcacagcacg cagcactag cgggcgcggt gccgtggagg aggtggatga ggagggcaag  
 1561 tttgtccggc tgcgcaacaa gtccaatgag gaccagtcca tgggcaattg gcagatcaag  
 1621 cgccagaatg gagatgatcc cttgctgact taccggttcc caccaaagtt caccctgaag  
 1681 gctgggcagg tggtgacgat ctgggctgca ggagctgggg ccacccacag cccccctacc  
 1741 gacctggtgt ggaaggcaca gaacacctgg ggctgcggga acagcctgcg tacggctctc  
 1801 atcaactcca ctggggaaga agtggccatg cgcaagctgg tgcgctcagt gactgtgggt  
 1861 gaggacgacg aggatgagga tggagatgac ctgctccatc accaccatgt gagtggtagc  
 1921 cgccgctgag gccgagcctg cactggggcc acccagccag gcctgggggc agcctctccc  
 1981 cagcctcccc gtgccaaaaa tcttttcatt aaagaatgtt tggaaacttt

Translation:

METPSQRRATRSGAQASSTPLSPTRITRLQEKEDLQELNDRLAVYIDRVRSLETENAG  
 LRLRITESEEVVSREVSGIKAAEYAEALGDARKTLDSVAKERARLQLELSKVREEFKEL  
 KARNTKKEGDLIAAQARLKDLEALLNSKEAALSTALSEKRTLEGELHDLRGQVAKLEA  
 ALGEAKKQLQDEMLRRVDAENRLQTMKEELDFQKNYSEELRETKRRHETRLVEIDNG  
 KQREFESRLADALQELRAQHEDQVEQYKKELEKTYSAKLDNARQSAERNNSNLVGAAHE  
 ELQQSRIRIDSLSAQLSQLQKQLAAKEAKLRDLEDSLARERDTSRLLAEKEREMAEM  
 RARMQQQLDEYQELLDIKLALDMEIHAYRKLLGEEERLRLSPSPSQRSGRGRASSHS  
 SQTQGGGSVTKKRKLESTESRSSFSQHARTSGRVAVEEVDEEGKFVRLRNKSNEQSM  
 GNWQIKRQNGDDPLLTYRFPFKETLKAGQVVTIWAAGAGATHSPPTDLVWKAQNTWGC  
 GNSLRTALINSTGEEVAMRKLVRSVTVVEDDEDEDGDDLLHHHHVSGSRR